

GenCore version 5.1.6
 OM protein - protein search, using sw model
 Run on: August 26, 2005, 19:04:47 ; Search time 171 Seconds
 (without alignments)
 726.024 Million cell updates/sec

Title: US-10-767-374-2
 Perfect score: 1688
 Sequence: 1 MGILLGLLLGLHVTYVGR.....AYIMLCRTKSQOEHHVEAAR 321
 Scoring table: BLOSUM62
 Gapop 10.0 , Gapext 0.5
 Searched: 2105692 seqs, 386760381 residues
 Total number of hits satisfying chosen parameters: 2105692
 Minimum DB seq length: 0
 Maximum DB seq length: 2000000000
 Post-processing: Minimum Match 0%
 Maximum Match 100%

Database : A_Geneseq_18Dec04.*
 Listing first 1500 summaries
 1: Geneseqp1980s.*
 2: Geneseqp1990s.*
 3: Geneseqp2000s.*
 4: Geneseqp2001s.*
 5: Geneseqp2002s.*
 6: Geneseqp2003as.*
 7: Geneseqp2003bs.*
 8: Geneseqp2004s.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

No.	Score	Match	Length	DB	ID	Description
RESULT 1						
ID	AA23322	standard; protein; 321 AA.				
DE	Amino acid sequence of the PRO362 polypeptide.					
PN	WO927098-A2.					
PD	03-JUN-1999.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 2;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 2						
ID	AA41691	standard; protein; 321 AA.				
DE	Human PRO 362 protein sequence.					
PN	WO9946281-A2.					
PD	16-SEP-1999.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 2;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 3						
ID	AAB33429	standard; protein; 321 AA.				
DE	Human PRO362 protein UNQ317 SEQ ID NO:80.					
PN	WO200053758-A2.					
PD	14-SEP-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 4						
ID	AA44247	standard; protein; 321 AA.				
DE	Human PRO362 (UNQ317) protein sequence SEQ ID NO:52.					
PN	WO200053756-A2.					
PD	14-SEP-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 5						
ID	AA95346	standard; protein; 321 AA.				
DE	Human PRO362 antitumour protein.					
PN	WO200037638-A2.					
PD	29-JUN-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 6						
ID	AA24047	standard; protein; 321 AA.				

DE Human PRO362 protein sequence SEQ ID NO:14.
 PN WO200053754-A1.
 PD 14-SEP-2000.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 3; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 7
 ID ABO25193 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050239-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 8
 ID ABU72199 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2002192706-A1.
 PD 19-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 9
 ID AB084879 standard; protein; 321 AA.
 DE Human secreted and transmembrane polypeptide PRO362.
 PN US2002177553-A1.
 PD 28-NOV-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 10
 ID ABU61077 standard; protein; 321 AA.
 DE Human PRO362 polypeptide.
 PN US2002169284-A1.
 PD 14-NOV-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 11
 ID ABU80346 standard; protein; 321 AA.
 DE Human secreted/transmembrane protein PRO362.
 PN US2003004102-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 12
 ID ABU07737 standard; protein; 321 AA.
 DE Human A-33 related antigen PRO362.
 PN US2002182206-A1.
 PD 05-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 13
 ID ADA24591 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050241-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 14
 ID ABO19648 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050240-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 15
 ID ADA12252 standard; protein; 321 AA.
 DE Human secreted/transmembrane polypeptide PRO362.

PN US2003055216-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 6; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 16
ID ABO19539 standard; protein; 321 AA.
DE Novel human secreted and transmembrane polypeptide #7.
PN US2003049633-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 6; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 17
ID ADB73558 standard; protein; 321 AA.
DE Human PRO polypeptide #7.
PN US2003045462-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 18
ID ADB76274 standard; protein; 321 AA.
DE Human PRO polypeptide #7.
PN US2003083248-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 19
ID ADC43700 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003054986-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 20
ID ADC61460 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003049684-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 21
ID ADC63424 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003054405-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 22
ID ADC66524 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003060406-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 23
ID ADC68648 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003064407-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 24
ID ADC62708 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003068648-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.

PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 25
ID ADC67773 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003069178-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 26
ID ADC41093 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003072745-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 27
ID ADC67148 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003073131-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 28
ID ADC62084 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003073624-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 29
ID ADC41717 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003104998-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 30
ID ADC78883 standard; protein; 321 AA.
DE Human PRO protein #56.
PN WO2003034984-A2.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 31
ID ADS49086 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003096744-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 32
ID ADE35140 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203434-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 33
ID ADE16254 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203435-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.

ID	ADB48386	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003104536-A1.				
PD	05-JUN-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 43					
ID	ADB89487	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003130181-A1.				
PD	10-JUL-2003.				
PA	(ASHK/) ASHKENAZI A J.				
PA	(BAKE/) BAKER K P.				
PA	(BOTS/) BOTSTEIN D.				
PA	(DESN/) DESNOVERS L.				
PA	(EATO/) EATON D L.				
PA	(FERR/) FERRARA N.				
PA	(FILV/) FILVAROFF E.				
PA	(FONG/) FONG S.				
PA	(GAOW/) GAO W.				
PA	(GERB/) GERBER H.				
PA	(GERR/) GERRITSEN M E.				
PA	(GODD/) GODDARD A.				
PA	(GODO/) GODOWSKI P J.				
PA	(GIRM/) GIRMALDI J C.				
PA	(GURN/) GURNEY A L.				
PA	(HILL/) HILLAN K J.				
PA	(KLJA/) KLJAVIN I J.				
PA	(KUOS/) KUO S S.				
PA	(NAPI/) NAPIER M A.				
PA	(PANJ/) PAN J.				
PA	(PAON/) PAONI N F.				
PA	(ROYM/) ROY M A.				
PA	(SHEL/) SHELTON D L.				
PA	(STEW/) STEWART T A.				
PA	(TUMA/) TUMAS D.				
PA	(WILL/) WILLIAMS P M.				
PA	(WOOD/) WOOD W I.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 44					
ID	ADF61127	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003195345-A1.				
PD	16-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 45					
ID	ADF39819	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003198994-A1.				
PD	23-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 46					
ID	ADF45615	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003195148-A1.				
PD	16-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 47					
ID	ADF24011	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003204055-A1.				
PD	30-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 48					
ID	ADF24011	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003204055-A1.				
PD	30-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 49					
ID	ADF24011	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				
DN	US2003204055-A1.				
PD	30-OCT-2003.				
PA	(GETH) GENENTECH INC.				
Query Match	100.0%;	Score 1688;	DB 8;	Length 321;	
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;			
RESULT 50					
ID	ADF24011	standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.				

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RESULT 48
ID ADF40443 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199021-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 49
ID ADF23387 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203402-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 50
ID ADF33370 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003194780-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 51
ID ADF26837 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199436-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 52
ID ADF27473 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199437-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 53
ID ADF41067 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199435-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 54
ID ADF32746 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003211091-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 55
ID ADF25112 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003211092-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 56
ID ADF26213 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199674-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 57
ID ADF32429 standard; protein; 321 AA.
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ID ADF34002 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003194410-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 58
ID ADF46239 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003195344-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 59
ID ADG50225 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003207803-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 60
ID ADG49601 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003215905-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 61
ID ADG51473 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003215908-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 62
ID ADG48977 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003216305-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 63
ID ADG48353 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003216560-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 64
ID ADG50849 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004005312-A1.
PD 08-JAN-2004.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 65
ID ADG58793 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004005657-A1.
PD 08-JAN-2004.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 66
ID ADG62249 standard; protein; 321 AA.
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DE Human secreted/transmembrane protein, PRO362.
PN US2004006219-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 67
ID ADH25274 standard; protein; 321 AA.
DE Human neurotrophin homologue related protein sequence SEQ ID NO:52.
PN EP1386931-A1.
PD 04-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 68
ID ADH17051 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004048332-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 69
ID ADL06885 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004063921-A1.
PD 01-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 70
ID ADN35285 standard; protein; 321 AA.
DE Human STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 71
ID AAB19396 standard; protein; 399 AA.
DE Amino acid sequence of a human secreted protein.
PN WO200061755-A2.
PD 19-OCT-2000.
PA (CHIR) CHIRON CORP.
Query Match 100.0%; Score 1688; DB 3; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 72
ID AAB04230 standard; protein; 399 AA.
DE Human gene 11 encoded secreted protein HMSOW51, SEQ ID NO:85.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 4; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 73
ID AAE04208 standard; protein; 399 AA.
DE Human gene 11 encoded secreted protein HMSOW51, SEQ ID NO:63.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 4; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 74
ID ABG64488 standard; protein; 399 AA.
DE Human albumin fusion protein #1163.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 5; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 75
ID ABG64487 standard; protein; 399 AA.
DE Human albumin fusion protein #1162.

PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 5; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 76
ID ABU03567 standard; protein; 399 AA.
DE Angiogenesis-associated human protein sequence #112.
PN WO200279492-A2.
PD 10-OCT-2002.
PA (EOSB-) EOS BIOTECHNOLOGY INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 77
ID ADA57183 standard; protein; 399 AA.
DE Human secreted protein #466.
PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 78
ID ADA41052 standard; protein; 399 AA.
DE Human secreted protein.
PN WO2002102993-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 79
ID ABR47904 standard; protein; 399 AA.
DE Human secreted protein, SEQ ID 795.
PN WO200295010-A2.
PD 28-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 80
ID ABR00157 standard; protein; 399 AA.
DE Human gene 147 encoded secreted protein HSDEK49, SEQ ID NO:446.
PN WO200276488-A1.
PD 03-OCT-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 81
ID ADB91646 standard; protein; 399 AA.
DE Human secreted protein #SEQ ID 592.
PN WO2003004622-A2.
PD 16-JAN-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 7; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 82
ID ADC74298 standard; protein; 399 AA.
DE Human secreted protein - SEQ ID 931.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 7; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 83
ID ADL77755 standard; protein; 399 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID NO 1237.
PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE/) ROSEN C A.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 84
ID ADL77754 standard; protein; 399 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID NO 1236.

PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE/) ROSEN C A.
PA (HASE/) HASELTINE W A.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 85
ID ADL67172 standard; protein; 399 AA.
DE Human B7-H6 protein SEQ ID NO:42.
PN WO2004022594-A2.
PD 18-MAR-2004.
PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 86
ID ADN35311 standard; protein; 399 AA.
DE Human STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 87
ID ABM2416 standard; protein; 399 AA.
DE Tumour-associated antigenic target (TAT) polypeptide PRO34043, SEQ:6205.
PN WO2004030615-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 88
ID ADP23762 standard; protein; 399 AA.
DE PRO polypeptide SEQ ID NO:940.
PN WO2004041170-A2.
PD 21-MAY-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 89
ID AAM3874 standard; protein; 399 AA.
DE Human polypeptide, SEQ ID NO: 3986.
PN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 99.5%; Score 1680; DB 4; Length 399;
Best Local Similarity 99.4%; Pred. No. 6.1e-135;
RESULT 90
ID ADL31953 standard; protein; 399 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3986.
PN EP1396543-A2.
PD 10-MAR-2004.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 99.5%; Score 1680; DB 8; Length 399;
Best Local Similarity 99.4%; Pred. No. 6.1e-135;
RESULT 91
ID AAE04290 standard; protein; 386 AA.
DE Human gene 11 encoded secreted protein fragment, SEQ ID NO:154.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 96.3%; Score 1625; DB 4; Length 386;
Best Local Similarity 100.0%; Pred. No. 3e-130;
RESULT 92
ID ADL67174 standard; protein; 281 AA.
DE Human B7-H6 (ECD) protein SEQ ID NO:44.
PN WO2004022594-A2.
PD 18-MAR-2004.
PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
Query Match 87.7%; Score 1480; DB 8; Length 281;
Best Local Similarity 100.0%; Pred. No. 4.9e-118;
RESULT 93
ID ADA57531 standard; protein; 305 AA.
DE Human secreted protein #466.

PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 94
ID ADA41415 standard; protein; 305 AA.
DE Human secreted protein.
PN WO2002102993-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 95
ID ABR48114 standard; protein; 305 AA.
DE Human secreted protein, SEQ ID 1005.
PN WO200295010-A2.
PD 28-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 96
ID ABR00282 standard; protein; 305 AA.
DE Human gene 147 encoded secreted protein HSDEK49, SEQ ID NO:571.
PN WO200276488-A1.
PD 03-OCT-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 97
ID ADB91804 standard; protein; 305 AA.
DE Human secreted protein #SEQ ID 750.
PN WO2003004622-A2.
PD 16-JAN-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 7; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 98
ID ADC74547 standard; protein; 305 AA.
DE Human secreted protein - SEQ ID 1180.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 7; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 99
ID ADN35312 standard; protein; 305 AA.
DE Human short STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 67.4%; Score 1137; DB 8; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 100
ID AAY30814 standard; protein; 306 AA.
DE Human secreted protein encoded from gene 4.
PN WO9940100-A1.
PD 12-AUG-1999.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 2; Length 306;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 101
ID AAM93588 standard; protein; 184 AA.
DE Human polypeptide, SEQ ID NO: 3387.
PN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 58.2%; Score 983; DB 4; Length 184;
Best Local Similarity 100.0%; Pred. No. 8.8e-76;
RESULT 102
ID ADL31354 standard; protein; 184 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3387.
PN EP1396543-A2.

PD 10-MAR-2004.
 PA (REAS-) RES ASSOC BIOTECHNOLOGY.
 Query Match 58.2%; Score 983; DB 8; Length 184;
 Best Local Similarity 100.0%; Pred. No. 8.8e-76;
 RESULT 103
 ID AAY82322 standard; protein; 175 AA.
 DE Human protein transport molecule (PTAM) SEQ ID NO:6.
 PD WO200012703-A2.
 PD 09-MAR-2000.
 PA (INCY-) INCYTE PHARM INC.
 Query Match 51.5%; Score 870; DB 3; Length 175;
 Best Local Similarity 64.7%; Pred. No. 3.8e-66;
 RESULT 104
 ID ADK70486 standard; protein; 175 AA.
 DE Respiratory disease differentially expressed protein #52.
 PD WO2003101283-A2.
 PD 11-DEC-2003.
 PA (INCY-) INCYTE CORP.
 Query Match 51.5%; Score 870; DB 8; Length 175;
 Best Local Similarity 64.7%; Pred. No. 3.8e-66;
 RESULT 105
 ID ADL67144 standard; protein; 280 AA.
 DE Mouse B7-H6 protein SEQ ID NO:14.
 PD WO2004022594-A2.
 PD 18-MAR-2004.
 PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
 Query Match 43.1%; Score 728; DB 8; Length 280;
 Best Local Similarity 47.0%; Pred. No. 1e-53;
 RESULT 106
 ID ADN35313 standard; protein; 280 AA.
 DE Murine STGMA protein.
 PD WO2004031105-A2.
 PD 15-APR-2004.
 PA (GETH-) GENENTECH INC.
 Query Match 43.1%; Score 728; DB 8; Length 280;
 Best Local Similarity 47.0%; Pred. No. 1e-53;
 RESULT 107
 ID ADL67146 standard; protein; 188 AA.
 DE Mouse B7-H6(ECD) protein SEQ ID NO:16.
 PD WO2004022594-A2.
 PD 18-MAR-2004.
 PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
 Query Match 36.3%; Score 613; DB 8; Length 188;
 Best Local Similarity 45.7%; Pred. No. 4e-44;
 RESULT 108
 ID AAE10596 standard; protein; 93 AA.
 DE Human macrophage-expressed protein #21.
 PD WO200164839-A2.
 PD 07-SEP-2001.
 PA (HYSE-) HYSEQ INC.
 Query Match 29.1%; Score 492; DB 4; Length 93;
 Best Local Similarity 98.9%; Pred. No. 3.5e-34;
 RESULT 109
 ID ADH80723 standard; protein; 300 AA.
 DE Human polypeptide #40.
 PD US2003232054-A1.
 PD 18-DEC-2003.
 PA (TANG/) TANG Y T.
 PA (LIUC/) LIU C.
 PA (ASUN/) ASUNDI V.
 PA (CHEN/) CHEN R.
 PA (QIAN/) QIAN X B.
 PA (WANG/) WANG Z W.
 PA (WEHR/) WEHRMAN T.
 PA (ZHAN/) ZHANG J.
 PA (ZHOU/) ZHOU P.
 PA (CAOY/) CAO Y.
 PA (DRMA/) DRMANAC R T.
 Query Match 10.7%; Score 180; DB 8; Length 300;
 Best Local Similarity 27.9%; Pred. No. 8.2e-07;
 RESULT 110
 ID AAY23321 standard; protein; 299 AA.
 DE Amino acid sequence of the PRO301 polypeptide.

PN WO9927098-A2.
 PD 03-JUN-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 111
 ID AAW74464 standard; protein; 299 AA.
 DE F11 antigen protein sequence.
 PD WO9902561-A1.
 PD 21-JAN-1999.
 PA (SMIK-) SMITHKLINE BEECHAM CORP.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 112
 ID AAY08071 standard; protein; 299 AA.
 DE Human PRO307 protein.
 PD WO9914241-A2.
 PD 25-MAR-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 113
 ID AAV13364 standard; protein; 299 AA.
 DE Amino acid sequence of protein PRO301.
 PD WO9914328-A2.
 PD 25-MAR-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 114
 ID AAY76011 standard; protein; 299 AA.
 DE Human A33 receptor homologue, SEQ ID NO:189.
 PD WO9955865-A1.
 PD 04-NOV-1999.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 115
 ID AAY76076 standard; protein; 299 AA.
 DE Human A33 receptor homologue, SEQ ID NO:331.
 PD WO9955865-A1.
 PD 04-NOV-1999.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 116
 ID AAY70670 standard; protein; 299 AA.
 DE Human PRO301 protein.
 PD WO200015797-A2.
 PD 23-MAR-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 117
 ID AAB24405 standard; protein; 299 AA.
 DE Human PRO301 protein sequence SEQ ID NO:90.
 PD WO200032221-A2.
 PD 08-JUN-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 118
 ID AAY95344 standard; protein; 299 AA.
 DE Human PRO301 antitumour protein.
 PD WO200037638-A2.
 PD 29-JUN-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 119
 ID AAB80232 standard; protein; 299 AA.
 DE Human PRO301 protein.
 PD WO200104311-A1.

PD 18-JAN-2001.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 120
ID AAM93577 standard; protein; 299 AA.
DE Human polypeptide, SEQ ID NO: 3365.
PN EPI130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 121
ID AAB56015 standard; protein; 299 AA.
DE Skin cell protein, SEQ ID NO: 331.
PN WO200069884-A2.
PD 23-NOV-2000.
PA (GENE-) GENESIS RES & DEV CORP LTD.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 122
ID AAB55950 standard; protein; 299 AA.
DE Skin cell protein, SEQ ID NO: 189.
PN WO200069884-A2.
PD 23-NOV-2000.
PA (GENE-) GENESIS RES & DEV CORP LTD.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 123
ID AAB31202 standard; protein; 299 AA.
DE Amino acid sequence of human polypeptide PRO301.
PN WO20007037-A2.
PD 21-DEC-2000.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 124
ID AAU00823 standard; protein; 299 AA.
DE Human immune response protein PRO301 (UNQ264).
PN WO200119991-A1.
PD 22-MAR-2001.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 125
ID AAU12354 standard; protein; 299 AA.
DE Human PRO301 polypeptide sequence.
PN WO200140466-A2.
PD 07-JUN-2001.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 126
ID AAB53086 standard; protein; 299 AA.
DE Human angiogenesis-associated protein PRO301, SEQ ID NO:119.
PN WO200053753-A2.
PD 14-SEP-2000.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 127
ID AAU14405 standard; protein; 299 AA.
DE Human novel protein #276.
PN WO200155437-A2.
PD 02-AUG-2001.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 128
ID AAU14404 standard; protein; 299 AA.
DE Human novel protein #275.
PN WO200155437-A2.
PD 02-AUG-2001.

PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 129
ID AAU14168 standard; protein; 299 AA.
DE Human novel protein #39.
PN WO200155437-A2.
PD 02-AUG-2001.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 130
ID AAE03896 standard; protein; 299 AA.
DE Human gene 23 encoded secreted protein fragment, SEQ ID NO:148.
PN WO200136440-A1.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 131
ID AAE03840 standard; protein; 299 AA.
DE Human gene 23 encoded secreted protein HACAA29, SEQ ID NO: 86.
PN WO200136440-A1.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 132
ID AAE03870 standard; protein; 299 AA.
DE Human gene 23 encoded secreted protein HACAA29, SEQ ID NO:116.
PN WO200136440-A1.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 133
ID ABB90290 standard; protein; 299 AA.
DE Human polypeptide SEQ ID NO 2666.
PN WO200190304-A2.
PD 29-NOV-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 134
ID ABB84843 standard; protein; 299 AA.
DE Human PRO301 protein sequence SEQ ID NO:54.
PN WO200200690-A2.
PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 135
ID ABG64551 standard; protein; 299 AA.
DE Human albumin fusion protein #1226.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 136
ID ABG64552 standard; protein; 299 AA.
DE Human albumin fusion protein #1227.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 137
ID ABB72215 standard; protein; 299 AA.
DE Human protein isolated from skin cells SEQ ID NO: 331.
PN WO200190357-A1.
PD 29-NOV-2001.
PA (GENE-) GENESIS RES & DEV CORP LTD.

Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 138
ID AB072150 standard; protein; 299 AA.
DE Human protein isolated from skin cells SEQ ID NO: 189.
PN W0200190357-A1.
PD 29-NOV-2001.
PA (GENE-) GENESIS RES & DEV CORP LTD.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 139
ID ABB95449 standard; protein; 299 AA.
DE Human angiogenesis related protein PRO301 SEQ ID NO: 54.
PN W0200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
PA (BAKE/) BAKER K P.
PA (FERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANU/) PAN J.
PA (PAON/) PAONI N F.
PA (STEP/) STEPHAN J F.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 140
ID AB071610 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2002146709-A1.
PD 10-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 141
ID ABO17798 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032156-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 142
ID AB071465 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2002192659-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 143
ID ABO25173 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003040014-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 144
ID AB081052 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003004311-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

RESULT 145
ID AB071911 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003003530-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 146
ID ABO01794 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2002197671-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 147
ID AB066752 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003036180-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 148
ID AB054367 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2002132240-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 149
ID AB067291 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032063-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 150
ID AB047382 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003044839-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 151
ID AB059833 standard; protein; 299 AA.
DE Novel secreted and transmembrane protein PRO301.
PN US2003017563-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 152
ID ABO25023 standard; protein; 299 AA.
DE Human secreted/transmembrane protein (PRO) #183.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 153
ID AB064519 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #23.
PN US2002160374-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 154
ID AB064519 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #23.
PN US2002160374-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

ID ABU72059 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2002177165-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 155
ID ABU67365 standard; protein; 299 AA.
DE Human secreted protein PRO301.
PN US2003023054-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 156
ID ABU67160 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032062-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 157
ID ABO14885 standard; protein; 299 AA.
DE Human secreted / transmembrane polypeptide PRO301.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 158
ID ABU07736 standard; protein; 299 AA.
DE Human A-33 related antigen PRO301.
PN US2002182206-A1.
PD 05-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 159
ID AAO16451 standard; protein; 299 AA.
DE Human junctional adhesion molecule 1 (hujami).
PN WO2003008541-A2.
PD 30-JAN-2003.
PA (ELIL) LILLY & CO ELI.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 160
ID ABU67028 standard; protein; 299 AA.
DE Human secreted/transmembrane, PRO, protein SEQ ID 366.
PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 161
ID ABU69642 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003017463-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 162
ID ABU79802 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003032057-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 163
ID ABO14824 standard; protein; 299 AA.
DE Human secreted / transmembrane polypeptide PRO301.
PN US2003027143-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 164
ID ADA45985 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 165
ID ADA76316 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 166
ID ADB29324 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003092002-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 167
ID ADA18966 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 168
ID ADA61589 standard; protein; 299 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 169
ID ADB19374 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 170
ID ADB27915 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 171
ID ADA86394 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 172
ID ADB15958 standard; protein; 299 AA.
DE Human PRO polypeptide #183.

PN US2003087350-A1.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 173
ID ADA47744 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 174
ID ADA18180 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039971-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 175
ID ABO32776 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003045693-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 176
ID ADA67539 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 177
ID ADB30546 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 178
ID ADA65842 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082693-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 179
ID ADA97054 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 180
ID ADA79358 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 181
ID ADA87497 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087345-A1.

PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 182
ID ADB16699 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 183
ID ABO34836 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003044793-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 184
ID ADA16155 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049621-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 185
ID ADA91791 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082694-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 186
ID ADB14854 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087351-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 187
ID ADA47263 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003044844-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 188
ID ADB18815 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 189
ID ADA94030 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 190
ID ADB19926 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082691-A1.
PD 01-MAY-2003.

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PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 191
ID ADB13238 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 192
ID ABO43331 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003044945-A1.
PD 06-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 193
ID ADA74492 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 194
ID ADA42300 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054401-A1.
PD 20-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 195
ID ADB24725 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077713-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 196
ID ADA82249 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 197
ID ADA75212 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 198
ID ADA85290 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 199
ID ADA84738 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 200
ID ABO17514 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003064367-A1.
PD 03-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 201
ID ADB29994 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 202
ID ADA80522 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 203
ID ADA75764 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 204
ID ADA46989 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 205
ID ADB25285 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077715-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 206
ID ADA93461 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077721-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 207
ID ADB26811 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 208
ID ADB31098 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
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Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 209
ID ADA61026 standard; protein; 299 AA.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 210
ID ADB24173 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077714-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 211
ID ADA96502 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082690-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 212
ID ADA81074 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 213
ID ADA95950 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082759-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 214
ID ADB26259 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 215
ID ADB21744 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 216
ID ADA77523 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068797-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 217
ID ADB18263 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US200307710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 218
ID ADA86946 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 219
ID ADA16579 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039969-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 220
ID ADA13008 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049622-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 221
ID ADA41876 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003082540-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 222
ID ADA88049 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 223
ID ADA46437 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 224
ID ADA17223 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003017498-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 225
ID ADA42726 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054351-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 226
ID ADB28467 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 227

ID ADB29019 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 228
ID ADA76971 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 229
ID ADA88601 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 230
ID ADA97606 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 231
ID ADB27363 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003022239-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 232
ID ADB22296 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087344-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 233
ID ABO19860 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO302.
PN US2003044902-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 234
ID ABO17575 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003064923-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 235
ID ADA66987 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 236
ID ADB22848 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077711-A1.
PD 24-APR-2003.

PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 237
ID ADB23621 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 238
ID ADA92343 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 239
ID ADB15406 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 240
ID ADB38658 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 241
ID ADB38106 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 242
ID ADB66578 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 243
ID ADB89658 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 244
ID ADB90390 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 245
ID ADB77645 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003077654-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 246
ID ADB39491 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 247
ID ADB74781 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003082542-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 248
ID ADB47114 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082687-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 249
ID ADB6721 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082697-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 250
ID ADB77326 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082696-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 251
ID ADB34483 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307717-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 252
ID ADB35587 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 253
ID ADB33931 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307716-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 254
ID ADB35035 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307718-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 255
ID ADB36139 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307720-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 256
ID ADB46534 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082692-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 257
ID ADC28427 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059772-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 258
ID ADC39627 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059828-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 259
ID ADC40141 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059829-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 260
ID ADC18969 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003036061-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 261
ID ADC34265 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003036094-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 262
ID ADC29320 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049676-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 263
ID ADC28851 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049677-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

RESULT 273

ID ADC60485 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
FN US2003087367-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 274

ID ADC50960 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
FN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 275

ID ADC65487 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
FN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 276

ID ADC54585 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
FN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 277

ID ADC53546 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
FN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 278

ID ADC59069 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
FN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 279

ID ADC55947 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
FN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 280

ID ADC58517 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
FN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 281

ID ADC12363 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
FN US2003082541-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 282

ID ADC12363 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
FN US2003082541-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

ID ADD03191 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092104-A1.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 283
ID ADC90183 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 284
ID ADC69602 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194770-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 285
ID ADC48491 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 286
ID ADD10020 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194776-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 287
ID ADD04595 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087334-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 288
ID ADC80551 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 289
ID ADD11058 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 290
ID ADD10343 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 291
ID ADC47939 standard; protein; 299 AA.

DE Human PRO polypeptide #183.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 292
ID ADD04918 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003104469-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 293
ID ADC79999 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 294
ID ADD11303 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 295
ID ADD09468 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 296
ID ADD03924 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003104381-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 297
ID ADD03500 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003108983-A1.
PD 12-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 298
ID ADD4181 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 299
ID ADD52320 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 300
ID ADD53060 standard; protein; 299 AA.
DE Human PRO polypeptide #183.

PN US2003194792-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 301
ID ADD53612 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203437-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 302
ID ADD37096 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 303
ID ADD51768 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 304
ID ADD02567 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 305
ID ADD02001 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 306
ID ADD54183 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 307
ID ADD92500 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 308
ID ADD91396 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 309
ID ADE04010 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199057-A1.

PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 310
ID ADE32307 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 311
ID ADE22239 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 312
ID ADD79463 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 313
ID ADE41999 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 314
ID ADE17816 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 315
ID ADD91948 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 316
ID ADE33411 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194767-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 317
ID ADE33963 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 318
ID ADD80015 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207417-A1.
PD 06-NOV-2003.

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PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 319
ID ADE42551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 320
ID ADE19472 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 321
ID ADE34752 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003077583-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 322
ID ADE18920 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 323
ID ADE43116 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 324
ID ADD95905 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 325
ID ADE22791 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199064-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 326
ID ADD78909 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203429-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 327
ID ADE32859 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 328
ID ADE42551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 329
ID ADD80567 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207418-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 330
ID ADD89595 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 331
ID ADE40879 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 332
ID ADE04678 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 333
ID ADE92807 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 334
ID ADG21516 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207355-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 335
ID ADG23157 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 336
ID ADF97492 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 337
ID ADE32859 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
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Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 337
ID ADG80556 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 338
ID ADG80004 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 339
ID ADG63772 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003170721-A1.
PD 11-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 340
ID ADH62528 standard; protein; 299 AA.
DE Human PRO301 protein.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUNA/) TUNAS D.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 341
ID ADH59235 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039972-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 342
ID ADH55296 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 343
ID ADH55848 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 344
ID ADH138014 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054352-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 345
ID ADI64067 standard; protein; 299 AA.

DE Novel human secreted and transmembrane protein PRO301.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 346
ID ADI65016 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 347
ID ADI63515 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 348
ID ADH81929 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 349
ID ADH81377 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 350
ID ADJ26282 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054349-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 351
ID ADM82546 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 352
ID ADM15945 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 353
ID ADNI6574 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 354
ID ADNI5393 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.

PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 355
ID ADN14841 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 356
ID ADC81103 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 357
ID ADE79197 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003135025-A1.
PD 17-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 358
ID ADD76551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 359
ID ADD87915 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 360
ID ADD86319 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 361
ID ADE79621 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003130489-A1.
PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 362
ID ADE75767 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 363
ID ADE73297 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003129592-A1.

PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 364
ID AD841304 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 365
ID ADE23343 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 366
ID ADE23895 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 367
ID ADE24538 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 368
ID ADD87363 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 369
ID AD889229 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 370
ID AD841186 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003104558-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 371
ID ADE73832 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148370-A1.
PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 372
ID ADE18368 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194794-A1.
PD 16-OCT-2003.

PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 373
ID ADE88677 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199054-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 374
ID ADE99386 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211576-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 375
ID ADE94697 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 376
ID ADE91108 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 377
ID ADE95249 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 378
ID ADE93359 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 379
ID ADF34940 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 380
ID ADE98505 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211569-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 381
ID ADE92255 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 382
ID ADE90556 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 383
ID ADE91703 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 384
ID ADE98932 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211568-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 385
ID ADG40402 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US200325253-A1.
PD 04-DEC-2003.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A L.
PA (MATH/) MATHER J P.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 386
ID ADF73796 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003180312-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 387
ID ADG02282 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 388
ID ADG22068 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 389
ID ADG20138 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 390
ID ADE92255 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

ID ADF98044 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 391
ID ADG24261 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 392
ID ADF98615 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 393
ID ADG03446 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 394
ID ADF99167 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 395
ID ADG16752 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 396
ID ADG05211 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 397
ID ADG19478 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 398
ID ADF73372 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003166051-A1.
PD 04-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 399
ID ADG13315 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 400
ID ADG08372 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 401
ID ADG15542 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 402
ID ADF96940 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 403
ID ADG06125 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 404
ID ADG23709 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 405
ID ADG03998 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 406
ID ADG24899 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207427-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 407
ID ADG07196 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 408
ID ADG07748 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.

PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 409
ID ADG55243 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 410
ID ADG60907 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 411
ID ADG62011 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 412
ID ADG92215 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003027145-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 413
ID ADG82212 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 414
ID ADG57451 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207362-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 415
ID ADG56899 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207364-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 416
ID ADG55795 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207365-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 417
ID ADG58555 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207368-A1.

PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 418
ID ADG70921 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207420-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 419
ID ADG92642 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003027146-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 420
ID ADG58003 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207363-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 421
ID ADG53587 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207415-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 422
ID ADG71473 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207421-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 423
ID ADG81660 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207805-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 424
ID ADH30622 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077723-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 425
ID ADG63621 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003180796-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 426
ID ADH11989 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207419-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 427
ID ADG52411 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207414-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 428
ID ADG54139 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207416-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 429
ID ADG81108 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194793-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 430
ID ADG56347 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207366-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 431
ID ADH21613 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207378-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 432
ID ADG61459 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207429-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 433
ID ADH28546 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003022331-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 434
ID ADG54691 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207367-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 435
ID ADG59731 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207369-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 436
ID ADH20431 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005553-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 437
ID ADH43487 standard; protein; 299 AA.
DE Human PRO polypeptide #27.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 438
ID ADH07286 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004006211-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 439
ID ADH59831 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003215904-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 440
ID ADH06859 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005665-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 441
ID ADI18155 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 442
ID ADI18601 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003152999-A1.
PD 14-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 443
ID ADI65321 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148419-A1.

PD 07-AUG-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 444
 ID ADI37584 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003096340-A1.
 PD 22-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 445
 ID ADG09898 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2004009548-A1.
 PD 15-JAN-2004.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 446
 ID ADH97380 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003190610-A1.
 PD 09-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 447
 ID ADI15369 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2003207382-A1.
 PD 06-NOV-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 448
 ID ADG09246 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2004009547-A1.
 PD 15-JAN-2004.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 449
 ID ADI65748 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003148371-A1.
 PD 07-AUG-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 450
 ID ADI14701 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2003207383-A1.
 PD 06-NOV-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 451
 ID ADH60491 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2004023331-A1.
 PD 05-FEB-2004.
 PA (DESN) DESNOYERS L.
 PA (GODD) GODDARD A.
 PA (GODO) GODOWSKI P J.
 PA (GUEN) GUERNEY A L.
 PA (MATH) MATHIER J P.
 PA (WILL) WILLIAMS P M.
 PA (WOOD) WOOD W I.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;

Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 452
 ID ADI18296 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2003207349-A1.
 PD 06-NOV-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 453
 ID ADJ99548 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003187238-A1.
 PD 02-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 454
 ID ADL08741 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003186358-A1.
 PD 02-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 455
 ID ADI47176 standard; protein; 299 AA.
 DE Human JAM-1 protein sequence.
 PN WO2004003145-A2.
 PD 08-JAN-2004.
 PA (NAST-) NASTECH PHARM CO INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 456
 ID ADM25082 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003096233-A1.
 PD 22-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 457
 ID ADK40844 standard; protein; 299 AA.
 DE Human platelet F11 receptor #1.
 PN US6699688-B1.
 PD 02-MAR-2004.
 PA (UJNY) UNIV NEW YORK STATE RES FOUND.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 458
 ID ADJ63577 standard; protein; 299 AA.
 DE Novel human secreted and transmembrane protein PRO301.
 PN US2004039164-A1.
 PD 26-FEB-2004.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 459
 ID ADM29832 standard; protein; 299 AA.
 DE Human secreted/transmembrane protein, #25.
 PN US2003190611-A1.
 PD 09-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 460
 ID ADL77818 standard; protein; 299 AA.
 DE Albumin fusion protein related therapeutic protein X, SEQ ID No 1300.
 PN US2004010134-A1.
 PD 15-JAN-2004.
 PA (ROSE) ROSEN C A.
 PA (HASE) HASELTINE W A.
 Query Match 10.6%; Score 178.5; DB 8; Length 299;

Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 461
ID ADL77819 standard; protein; 299 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID No 1301.
PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE//) ROSEN C A.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 462
ID ADJ77472 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004038336-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 463
ID ADK82832 standard; protein; 299 AA.
DE Human PRO polypeptide #27.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 464
ID ADJ65594 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004038335-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 465
ID ADL13132 standard; protein; 299 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3365.
PN EPI396543-A2.
PD 10-MAR-2004.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 466
ID ADM27730 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004048333-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 467
ID ADL26800 standard; protein; 299 AA.
DE Human JAM1 protein SEQ ID NO:54.
PN WO2004022778-A1.
PD 18-MAR-2004.
PA (GARV-) GARVAN INST MEDICAL RES.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 468
ID ADM42454 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004058424-A1.
PD 25-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 469
ID ADO06154 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US6686451-B1.
PD 03-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 470
ID ADN35284 standard; protein; 299 AA.
DE Human PRO301 protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 471
ID ADN05140 standard; protein; 299 AA.
DE Antipsooriatic protein sequence #749.
PN WO2004028479-A2.
PD 08-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 472
ID ADM28316 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004077064-A1.
PD 22-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 473
ID ADQ95890 standard; protein; 299 AA.
DE T cell activation associated protein #34.
PN WO2004058805-A2.
PD 15-JUL-2004.
PA (ASAH-) ASahi KASEI PHARMA CORP.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 474
ID ADL11006 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004137561-A1.
PD 15-JUL-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 475
ID ADL17915 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004147017-A1.
PD 29-JUL-2004.
PA (ASHK/) ASHKENAZI A.
PA (BOTS/) BOTSTEIN D.
PA (DESN/) DESNOYERS L.
PA (EATO/) EATON D L.
PA (FERR/) FERRARA N.
PA (FILV/) FILVAROFF E.
PA (FONG/) FONG S.
PA (GAOW/) GAO W.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GRIM/) GRIMALDI C J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (KLJA/) KLJAVIN I J.
PA (MATH/) MATHER J P.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (ROIM/) ROY M A.
PA (STEW/) STEWART T A.
PA (TUNA/) TUMAS D.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 476

ID ADR27641 standard; protein; 299 AA.
DE Human F11 receptor protein Seq 7.
PN WO2004063327-A2.
PD 29-JUL-2004.
PA (KORN//) KORNECKI E.
PA (BABI//) BABINSKA A.
PA (EHLR//) EHRLICH Y H.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 477
ID ADI95798 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077659-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 478
ID ADI96350 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207354-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 479
ID ADR46577 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4, SEQ ID 8.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 480
ID ADR46571 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 481
ID ADR46573 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 482
ID ADR46579 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 5.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 483
ID ADT94260 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003152922-A1.
PD 14-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 484
ID ADT94260 standard; protein; 299 AA.
DE Human PRO301 protein.
PN AU2003259607-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

RESULT 485
ID ADS74554 standard; protein; 299 AA.
DE Human secreted/transmembrane protein #25.
PN US2004185531-A1.
PD 23-SEP-2004.
PA (ASHK/) ASHKENAZI A.
PA (BOTS/) BOTSTEIN D.
PA (DESN/) DESNOYERS L.
PA (EATO/) EATON D L.
PA (FERR/) FERRARA N.
PA (FILV/) FILVAROFF E.
PA (FONG/) FONG S.
PA (GAOW/) GAO W.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GRIM/) GRIMALDI C J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (KLJA/) KLJAVIN I J.
PA (MATH/) MATHER J P.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (ROYM/) ROY M A.
PA (STEW/) STEWART T A.
PA (TUMA/) TUMAS D.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 486
ID ADS09073 standard; protein; 320 AA.
DE Novel protein-related contig polypeptide sequence #139.
PN WO2003054152-A2.
PD 03-JUL-2003.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 7; Length 320;
Best Local Similarity 26.5%; Pred. No. 1.2e-06;
RESULT 487
ID ADU67616 standard; protein; 335 AA.
DE Human ovarian specific polypeptide SEQ ID NO:330.
PN WO2004013311-A2.
PD 12-FEB-2004.
PA (DIAD-) DIADEXUS INC.
Query Match 10.6%; Score 178.5; DB 8; Length 335;
Best Local Similarity 26.5%; Pred. No. 1.3e-06;
RESULT 488
ID ADR08038 standard; protein; 336 AA.
DE Novel protein (useful for identifying genetic disorders) #193.
PN WO2003054152-A2.
PD 03-JUL-2003.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 7; Length 336;
Best Local Similarity 26.5%; Pred. No. 1.3e-06;
RESULT 489
ID AAW14146 standard; protein; 319 AA.
DE Human A33 antigen.
PN WO9708189-A1.
PD 06-MAR-1997.
PA (LUDW-) LUDWIG INST CANCER RES.
Query Match 10.5%; Score 177; DB 2; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 490
ID AAT23323 standard; protein; 319 AA.
DE Amino acid sequence of the A33 antigen.
PN WO9927098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.5%; Score 177; DB 2; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 491
ID AAB65863 standard; protein; 319 AA.

DE Human A33 protein SEQ ID NO: 67.
 PN WO200078808-A1.
 PA (MILL-) MILLENNIUM PHARM INC.
 Query Match 10.5%; Score 177; DB 4; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 492
 ID ADA10947 standard; protein; 319 AA.
 DE Human cDNA differentially expressed in colon cancer #43 product.
 PN US2002160382-A1.
 PD 31-OCT-2002.
 PA (LASE/) LASEK A W.
 PA (JONE/) JONES D A.
 Query Match 10.5%; Score 177; DB 6; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 493
 ID ADH62533 standard; protein; 319 AA.
 DE Human A33 antigenic protein.
 PN US2003171568-A1.
 PD 11-SEP-2003.
 PA (ASHK/) ASHKENAZI A.
 PA (FONG/) FONG S.
 PA (GODD/) GODDARD A.
 PA (GURN/) GURNEY A L.
 PA (NAPI/) NAPIER M A.
 PA (TUMA/) TUNAS D.
 PA (WOOD/) WOOD W I.
 Query Match 10.5%; Score 177; DB 7; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 494
 ID ADN39847 standard; protein; 319 AA.
 DE Cancer/angiogenesis/fibrosis-related polypeptide, SEQ ID NO:C217.
 PN WO2003042661-A2.
 PD 22-MAY-2003.
 PA (EOSB-) EOS BIOTECHNOLOGY INC.
 Query Match 10.5%; Score 177; DB 7; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 495
 ID ADN35289 standard; protein; 319 AA.
 DE Human A33 antigen protein.
 PN WO2004031105-A2.
 PD 15-APR-2004.
 PA (GETH) GENENTECH INC.
 Query Match 10.5%; Score 177; DB 8; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 496
 ID ADP54587 standard; protein; 319 AA.
 DE Human PRO protein sequence SEQ ID NO:563.
 PN WO2004039956-A2.
 PD 13-MAY-2004.
 PA (GETH) GENENTECH INC.
 Query Match 10.5%; Score 177; DB 8; Length 319;
 Best Local Similarity 27.5%; Pred. No. 1.6e-06;
 RESULT 497
 ID ABP62881 standard; protein; 336 AA.
 DE Human polypeptide SEQ ID NO 318.
 PN WO200218424-A2.
 PD 07-MAR-2002.
 PA (HYSE-) HYSEQ INC.
 Query Match 10.5%; Score 177; DB 5; Length 336;
 Best Local Similarity 27.5%; Pred. No. 1.7e-06;
 RESULT 498
 ID ADC78439 standard; protein; 299 AA.
 DE Human PRO301 protein.
 PN WO200015796-A2.
 PD 23-MAR-2000.
 PA (GETH) GENENTECH INC.
 Query Match 10.5%; Score 176.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.6e-06;
 RESULT 499
 ID AAY08073 standard; protein; 268 AA.
 DE Human A33 protein.
 PN WO9914241-A2.

PD 25-MAR-1999.
 PA (GETH) GENENTECH INC.
 Query Match 10.4%; Score 176; DB 2; Length 268;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 500
 ID ADH62551 standard; protein; 268 AA.
 DE Human A33 antigenic protein fragment #1.
 PN US2003171568-A1.
 PD 11-SEP-2003.
 PA (ASHK/) ASHKENAZI A.
 PA (FONG/) FONG S.
 PA (GODD/) GODDARD A.
 PA (GURN/) GURNEY A L.
 PA (NAPI/) NAPIER M A.
 PA (TUMA/) TUNAS D.
 PA (WOOD/) WOOD W I.
 Query Match 10.4%; Score 176; DB 7; Length 268;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 501
 ID AAY23327 standard; protein; 270 AA.
 DE An A33 related antigen sequence.
 PN WO9927098-A2.
 PD 03-JUN-1999.
 PA (GETH) GENENTECH INC.
 Query Match 10.4%; Score 176; DB 2; Length 270;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 502
 ID AAY23329 standard; protein; 273 AA.
 DE An A33 related antigen sequence.
 PN WO9927098-A2.
 PD 03-JUN-1999.
 PA (GETH) GENENTECH INC.
 Query Match 10.4%; Score 176; DB 2; Length 273;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 503
 ID AAY08075 standard; protein; 273 AA.
 DE Human A33 protein fragment #2.
 PN WO9914241-A2.
 PD 25-MAR-1999.
 PA (GETH) GENENTECH INC.
 Query Match 10.4%; Score 176; DB 2; Length 273;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 504
 ID ADH62553 standard; protein; 273 AA.
 DE Human A33 antigenic protein fragment #2.
 PN US2003171568-A1.
 PD 11-SEP-2003.
 PA (ASHK/) ASHKENAZI A.
 PA (FONG/) FONG S.
 PA (GODD/) GODDARD A.
 PA (GURN/) GURNEY A L.
 PA (NAPI/) NAPIER M A.
 PA (TUMA/) TUNAS D.
 PA (WOOD/) WOOD W I.
 Query Match 10.4%; Score 176; DB 7; Length 273;
 Best Local Similarity 27.7%; Pred. No. 1.6e-06;
 RESULT 505
 ID ADH80722 standard; protein; 301 AA.
 DE Human polypeptide #39.
 PN US2003232054-A1.
 PD 18-DEC-2003.
 PA (TANG/) TANG Y T.
 PA (LIUC/) LIU C.
 PA (ASUN/) ASUNDI V.
 PA (CHEN/) CHEN R.
 PA (QIAN/) QIAN X B.
 PA (WEHR/) WEHRMAN T.
 PA (ZHAN/) ZHANG J.
 PA (ZHOU/) ZHOU P.
 PA (CAOI/) CAO Y.
 PA (DRMA/) DRMANAC R T.
 Query Match 10.3%; Score 173.5; DB 8; Length 301;

Best Local Similarity 26.6%; Pred. No. 3e-06;
RESULT 506
ID ADK40854 standard; protein; 316 AA.
DE Human A33 molecule.
PN US6699688-B1.
PD 02-MAR-2004.
PA (UUNY) UNIV NEW YORK STATE RES FOUND.
Query Match 10.2%; Score 172.5; DB 8; Length 316;
Best Local Similarity 26.4%; Pred. No. 3.9e-06;
RESULT 507
ID ADQ89964 standard; protein; 34350 AA.
DE Antagonist of cell cycle progression polypeptide #197.
PN WO2004063362-A2.
PD 29-JUL-2004.
PA (CYCL-) CYCLACEL LTD.
Query Match 10.2%; Score 172; DB 8; Length 34350;
Best Local Similarity 27.5%; Pred. No. 0.0024;
RESULT 508
ID AAY23326 standard; protein; 260 AA.
DE An A33 related antigen sequence.
PN WO9227098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 509
ID AAY08072 standard; protein; 260 AA.
DE Human DNA40628 protein.
PN WO9914241-A2.
PD 25-MAR-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 510
ID ADHG2550 standard; protein; 260 AA.
DE Human PRO301 protein fragment #1.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.0%; Score 169.5; DB 7; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 511
ID AAY23328 standard; protein; 263 AA.
DE An A33 related antigen sequence.
PN WO9227098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.5e-06;
RESULT 512
ID AAY08074 standard; protein; 263 AA.
DE Human DNA40628 protein fragment #2.
PN WO9914241-A2.
PD 25-MAR-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.5e-06;
RESULT 513
ID ADHG2552 standard; protein; 263 AA.
DE Human PRO301 protein fragment #2.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 9.9%; Score 167; DB 7; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 519
ID ADK40853 standard; protein; 300 AA.
DE Mouse junction adhesion molecule (JAM).
PN US6699688-B1.
PD 02-MAR-2004.
PA (UUNY) UNIV NEW YORK STATE RES FOUND.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 520
ID ADN35293 standard; protein; 300 AA.
DE Human JAM protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 521
ID ABB83928 standard; protein; 365 AA.
DE PCAR SEQ ID NO 4.
PN US2002059654-A1.
PD 16-MAY-2002.
PA (BUHL/) BUHLER T.
PA (GADI/) GADIENT R A.

PA (TUMA/) TUMAS D.
Query Match 10.0%; Score 169.5; DB 7; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.5e-06;
RESULT 514
ID ADK46581 standard; protein; 300 AA.
DE Mouse junctional adhesion molecule-1, SEQ ID 12.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKO-) DOKURITSU GYOSEI HOJIN KAKAKU GIJUTSU SH.
Query Match 10.0%; Score 168; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 8.8e-06;
RESULT 515
ID AAW61379 standard; protein; 298 AA.
DE Human junctional adhesion molecule protein.
PN WO9824897-A1.
PD 11-JUN-1998.
PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
Query Match 9.9%; Score 167; DB 2; Length 298;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 516
ID AAW61380 standard; protein; 300 AA.
DE Mouse junctional adhesion molecule protein.
PN WO9824897-A1.
PD 11-JUN-1998.
PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
Query Match 9.9%; Score 167; DB 2; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 517
ID AAY23325 standard; protein; 300 AA.
DE A33 related antigen JAM.
PN WO9227098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 9.9%; Score 167; DB 2; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 518
ID ADHG2537 standard; protein; 300 AA.
DE Murine JAM protein used in the exemplification of the invention.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 9.9%; Score 167; DB 7; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 519
ID ADK40853 standard; protein; 300 AA.
DE Mouse junction adhesion molecule (JAM).
PN US6699688-B1.
PD 02-MAR-2004.
PA (UUNY) UNIV NEW YORK STATE RES FOUND.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 520
ID ADN35293 standard; protein; 300 AA.
DE Human JAM protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 521
ID ABB83928 standard; protein; 365 AA.
DE PCAR SEQ ID NO 4.
PN US2002059654-A1.
PD 16-MAY-2002.
PA (BUHL/) BUHLER T.
PA (GADI/) GADIENT R A.

PA (KORN/) KORN R.
PA (MOVV/) MOVVA R.
Query Match 9.8%; Score 166; DB 5; Length 365;
Best Local Similarity 24.2%; Pred. No. 1.7e-05;
RESULT 522
ID ABB83927 standard; protein; 261 AA.
DE C-terminally truncated pCAR SEQ ID NO 2.
PN US2002059654-A1.
PD 16-MAY-2002.
PA (BUHL/) BUHLER T.
PA (GADI/) GADIENT R. A.
PA (KORN/) KORN R.
PA (MOVV/) MOVVA R.
Query Match 9.7%; Score 163.5; DB 5; Length 261;
Best Local Similarity 27.7%; Pred. No. 1.8e-05;
RESULT 523
ID AAW14158 standard; protein; 318 AA.
DE Mouse A33 antigen.
PN WO9708189-A1.
PD 06-MAR-1997.
PA (LUDW-) LUDWIG INST CANCER RES.
Query Match 9.6%; Score 162.5; DB 2; Length 318;
Best Local Similarity 22.9%; Pred. No. 2.8e-05;
RESULT 524
ID AAB33253 standard; protein; 280 AA.
DE Gene 15 human secreted protein homologous amino acid sequence #133.
PN WO2000056754-A1.
PD 28-SEP-2000.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 9.5%; Score 160; DB 3; Length 280;
Best Local Similarity 24.7%; Pred. No. 3.9e-05;
RESULT 525
ID ADA50171 standard; protein; 412 AA.
DE Human CAR/mouse SCF fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOY/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 9.4%; Score 158.5; DB 6; Length 412;
Best Local Similarity 24.8%; Pred. No. 8.7e-05;
RESULT 526
ID AAY72878 standard; protein; 352 AA.
DE Human PRO5723 protein encoded by DNA82361 cDNA clone.
PN WO200116319-A2.
PD 08-MAR-2001.
PA (GETH-) GENENTECH INC.
Query Match 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 527
ID AAB50930 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN WO2000073452-A2.
PD 07-DEC-2000.
PA (GETH-) GENENTECH INC.
Query Match 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 528
ID AAB65294 standard; protein; 352 AA.
DE Human PRO5723 protein sequence SEQ ID NO:505.
PN WO2000073454-A1.
PD 07-DEC-2000.
PA (GETH-) GENENTECH INC.
Query Match 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 529
ID ABB84956 standard; protein; 352 AA.
DE Human PRO5723 protein sequence SEQ ID NO:280.
PN WO200200690-A2.
PD 03-JAN-2002.
PA (GETH-) GENENTECH INC.
Query Match 9.4%; Score 158; DB 5; Length 352;

Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 530
ID ABB95562 standard; protein; 352 AA.
DE Human angiogenesis related protein PRO5723 SEQ ID NO: 280.
FN WO200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
PA (BAKE/) BAKER K P.
PA (PERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N P.
PA (STEP/) STEPHAN J P.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 9.4%; Score 158; DB 5; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 531
ID ABU58109 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
FN US2003027163-A1.
PD 06-FEB-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 532
ID ABU59187 standard; protein; 352 AA.
DE Novel human secreted or transmembrane protein PRO5723.
FN US2002132252-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 533
ID ABU82699 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
FN US2003032023-A1.
PD 13-FEB-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 534
ID ABU60618 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
FN US2002160384-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 535
ID ABU80846 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
FN US2003036635-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 536
ID ABO33912 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
FN US2003045687-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 537
ID ABU14000 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
FN US2003045687-A1.

PN US2002103125-A1.
PD 01-AUG-2002.
PA (GETH) GENENTECH LTD. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 538
ID ABU72595 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200303531-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 539
ID ABG74762 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2002192752-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 540
ID ABU59334 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
PN US2003027162-A1.
PD 06-FEB-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 541
ID ABO26031 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2002127576-A1.
PD 12-SEP-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 542
ID ABU82155 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308063-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 543
ID ABU59040 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 544
ID ABU92418 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003022187-A1.
PD 30-JAN-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 545
ID ABU59483 standard; protein; 352 AA.
DE Novel human secreted or transmembrane protein PRO3301.
PN US2003027985-A1.
PD 06-FEB-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 546
ID ABU92249 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 547
ID ABU92249 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;

RESULT 547
ID ABU10955 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2002123463-A1.
PD 05-SEP-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 548
ID ABU81707 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002177164-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 549
ID ABU8646 standard; protein; 352 AA.
DE Human secreted and transmembrane polypeptide PRO5723.
PN US2002197615-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 550
ID ABO34160 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003017981-A1.
PD 23-JAN-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 551
ID ABJ72335 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2003050448-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 552
ID ADA38016 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003008297-A1.
PD 09-JAN-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 553
ID ADA21702 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO5723.
PN US2003054404-A1.
PD 20-MAR-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 554
ID ADA10489 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, PRO5723.
PN US2003059831-A1.
PD 27-MAR-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 555
ID ADA18033 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003054987-A1.
PD 20-MAR-2003.
Query Match Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 556
ID ADA28141 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003054359-A1.
PD 20-MAR-2003.

Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 557
ID ADA94721 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059832-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 558
ID ADA38946 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059780-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 559
ID ABJ72463 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2003027988-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 560
ID ADA93067 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003060407-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 561
ID ABO34358 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO 5723.
PN US2003044934-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 562
ID ABO53246 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003044806-A1.
PD 06-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 563
ID ADA22628 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO5723.
PN US2003040473-A1.
PD 27-FEB-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 564
ID ABO22616 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003017982-A1.
PD 23-JAN-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 565
ID ADA06794 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #141.
PN US2003049638-A1.
PD 13-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 566
ID ABJ72165 standard; protein; 352 AA.
DE Human membrane bound receptor/protein PRO5723 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.

Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 567
ID ADA39487 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059782-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 568
ID ADB83706 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073814-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 569
ID ADB80812 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308068-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 570
ID ADB73353 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096968-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 571
ID ADB96513 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003054403-A1.
PD 20-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 572
ID ADB78435 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092889-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 573
ID ADB85083 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003073817-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 574
ID ADB78189 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092886-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 575
ID ADB87255 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US200308067-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 576
ID ADB72165 standard; protein; 352 AA.
DE Human membrane bound receptor/protein PRO5723 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.

ID ADB84837 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003092890-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 577
ID ADB83952 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003069397-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 578
ID ADB73107 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092887-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 579
ID ADC57985 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003027754-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 580
ID ADC55349 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003045463-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 581
ID ADC12216 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003049681-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 582
ID ADC56638 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003064375-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 583
ID ADC11683 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003069403-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 584
ID ADC36945 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003088065-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 585
ID ADC21935 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096969-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 586
ID ADC49966 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088064-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 587
ID ADC49165 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088070-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 588
ID ADC49682 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 589
ID ADC47543 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088072-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 590
ID ADC14805 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003082546-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 591
ID ADC47288 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105288-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 592
ID ADD08337 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003068623-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 593
ID ADC82162 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003083461-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 594
ID ADD07804 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002193299-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 595
ID ADC78163 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.

PN US2003096972-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 596
ID ADC82695 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003059833-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 597
ID ADD06398 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073816-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 598
ID ADD10569 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 599
ID ADD08875 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073090-A1.
PD 17-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 600
ID ADC77917 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308066-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 601
ID ADD07124 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002193300-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 602
ID ADD11529 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 603
ID ADC83371 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003059783-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 604
ID ADD50880 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105291-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 605
ID ADD15380 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003059437-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 606
ID ADD51126 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105290-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 607
ID ADD55478 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003077593-A1.
PD 24-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 608
ID ADD37322 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 609
ID ADD56436 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003077594-A1.
PD 24-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 610
ID ADD50607 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096971-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 611
ID ADD54874 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2002132253-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 612
ID ADD50361 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096970-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 613
ID ADD51372 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105289-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 614
ID ADE31893 standard; protein; 352 AA.

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DE Human secreted/transmembrane protein PRO5723.
PN US2003068647-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 615
ID ADE27028 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003087304-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 616
ID ADE26495 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003087305-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 617
ID ADE67432 standard; protein; 352 AA.
DE Human PRO5723 amino acid sequence SEQ ID NO:505.
PN US2002198148-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 618
ID ADE94085 standard; protein; 352 AA.
DE Immune disease treatment/diagnosis related PRO5723.
PN US2003082199-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 619
ID ADI35686 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003050457-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 620
ID ADI00179 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003049682-A1.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 621
ID ADC48919 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092888-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 622
ID ADE21090 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100735-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 623
ID ADE05934 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100728-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 624
ID ADD75163 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100712-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 625
ID ADD75909 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100717-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 626
ID ADD85141 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100722-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 627
ID ADD86967 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100738-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 628
ID ADE20844 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100734-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 629
ID ADE39141 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096362-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 630
ID ADE05688 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100727-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 631
ID ADD73673 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100711-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 632
ID ADD78513 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100737-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 633
ID ADD78513 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100737-A1.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
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ID ADE41530 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 634
ID ADE21336 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100736-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 635
ID ADD77451 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100732-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 636
ID ADE20598 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100733-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 637
ID ADD75663 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100064-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 638
ID ADD74179 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100708-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 639
ID ADD74425 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100709-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 640
ID ADD76155 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100718-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 641
ID ADD85647 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100721-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 642
ID ADE05196 standard; protein; 352 AA.

DE Human PRO polypeptide #108.
PN US2003100726-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 643
ID ADD75409 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100714-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 644
ID ADD76953 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100715-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 645
ID ADD86721 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100719-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 646
ID ADD78189 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100731-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 647
ID ADD77697 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100729-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 648
ID ADD77943 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100730-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 649
ID ADD85401 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100725-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 650
ID ADD73933 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100710-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 651
ID ADD74671 standard; protein; 352 AA.
DE Human PRO polypeptide #108.

PN US2003100713-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 652
ID ADG12125 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100716-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 653
ID ADG85893 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100720-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 654
ID ADE05442 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100723-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 655
ID ADD74917 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100724-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 656
ID ADF35631 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003194760-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 657
ID ADG11881 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003228655-A1.
PD 11-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 658
ID ADG05729 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096959-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 659
ID ADG27283 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096962-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 660
ID ADG11346 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096967-A1.
PD 22-MAY-2003.

PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 661
ID ADG12125 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096963-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 662
ID ADF94682 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096964-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 663
ID ADG06778 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096966-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 664
ID ADH39122 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096965-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 665
ID ADH19751 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003228656-A1.
PD 11-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 666
ID ADH21244 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003224358-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 667
ID ADH20284 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003219856-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 668
ID ADH43713 standard; protein; 352 AA.
DE Human PRO polypeptide #140.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 669
ID ADG34212 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2004006206-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;

Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 670
ID ADI33682 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096960-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 671
ID ADH69776 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2004019183-A1.
PD 29-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 672
ID ADI29937 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096961-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 673
ID ADM27334 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200404179-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 674
ID ADK83058 standard; protein; 352 AA.
DE Human PRO polypeptide #140.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 675
ID ADK66692 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US200404180-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 676
ID AAW69697 standard; protein; 365 AA.
DE Human coxsackievirus and Ad2 and Ad5 receptor HCAR protein.
PN WO9833819-A1.
PD 06-AUG-1998.
PA (UUNY) UNIV NEW YORK STATE.
Query Match 9.4%; Score 158; DB 2; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 677
ID AAW57212 standard; protein; 365 AA.
DE Human coxsackievirus and adenovirus receptor.
PN WO9811221-A2.
PD 19-MAR-1998.
PA (DAND) DANA FARBER CANCER INST INC.
Query Match 9.4%; Score 158; DB 2; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 678
ID AAB47270 standard; protein; 365 AA.
DE Human CAR.
PN US6245966-B1.
PD 12-JUN-2001.
PA (UYTE-) UNIV TECHNOLOGY CORP.
Query Match 9.4%; Score 158; DB 4; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;

RESULT 679
ID ABB08040 standard; protein; 365 AA.
DE Human coxsackie-adenovirus receptor (CAR).
PN WO200229072-A2.
PD 11-APR-2002.
PA (NOVS) NOVARTIS AG.
PA (NOVS) NOVARTIS-ERFINDUNGEN VERW GES MBH.
Query Match 9.4%; Score 158; DB 5; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 680
ID ABJ37063 standard; protein; 365 AA.
DE Human breast cancer / ovarian cancer related protein #39.
PN WO2003000012-A2.
PD 03-JAN-2003.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 9.4%; Score 158; DB 6; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 681
ID AD897544 standard; protein; 365 AA.
DE Human CAR wild-type protein.
PN WO2003070915-A2.
PD 28-AUG-2003.
PA (USSH) US DEPT HEALTH & HUMAN SERVICES.
Query Match 9.4%; Score 158; DB 7; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 682
ID ADN95226 standard; protein; 365 AA.
DE Human BEC/LEC-related protein sequence SeqID148.
PN WO2003080640-A1.
PD 02-OCT-2003.
PA (LUDW-) LUDWIG INST CANCER RES.
PA (LICN) LICENTIA LTD.
Query Match 9.4%; Score 158; DB 7; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 683
ID ABUL2046 standard; protein; 505 AA.
DE Human NOV4a CG59871-01 protein SEQ ID 12.
PN WO200281625-A2.
PD 17-OCT-2002.
PA (CURA-) CURAGEN CORP.
Query Match 9.4%; Score 158; DB 6; Length 505;
Best Local Similarity 23.3%; Pred. No. 0.00013;
RESULT 684
ID ADJ67617 standard; protein; 351 AA.
DE Human ovarian specific polypeptide SEQ ID NO:331.
PN WO2004013111-A2.
PD 12-FEB-2004.
PA (DIAD-) DIADEXUS INC.
Query Match 9.3%; Score 157.5; DB 8; Length 351;
Best Local Similarity 23.0%; Pred. No. 8.6e-05;
RESULT 685
ID ADA50172 standard; protein; 493 AA.
DE Human CAR/mouse anti-CD34 antibody fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 9.3%; Score 157.5; DB 6; Length 493;
Best Local Similarity 23.7%; Pred. No. 0.00014;
RESULT 686
ID AAW82729 standard; protein; 264 AA.
DE Adenovirus pACTSG2-SCAR protein.
PN WO9854346-A1.
PD 03-DEC-1998.
PA (GENV-) GENVEC INC.
Query Match 9.3%; Score 156.5; DB 2; Length 264;
Best Local Similarity 25.9%; Pred. No. 7.1e-05;
RESULT 687
ID AAW82730 standard; protein; 277 AA.
DE Adenovirus SCAR.RGD protein.
PN WO9854346-A1.

PD 03-DEC-1998.
 PA (GENV-) GENVEC INC. 9.3%; Score 156.5; DB 2; Length 277;
 Best Local Similarity 25.9%; Pred. No. 7.6e-05;
 RESULT 688
 ID AAU83699 standard; protein; 290 AA.
 DE Human PRO protein, Seq ID No 216.
 PN WO200208288-A2.
 PD 31-JAN-2002.
 PA (GETH) GENENTECH INC.
 Query Match
 Best Local Similarity 25.9%; Score 156.5; DB 5; Length 290;
 RESULT 689
 ID AAU82731 standard; protein; 397 AA.
 DE Adenovirus PACSG2SCAR.sig chimeric protein.
 PN WO9854346-A1.
 PD 03-DEC-1998.
 PA (GENV-) GENVEC INC. 9.3%; Score 156.5; DB 2; Length 397;
 Best Local Similarity 25.9%; Pred. No. 0.00012;
 RESULT 690
 ID AD450170 standard; protein; 412 AA.
 DE Human CAR/SCF fusion protein.
 PN US2003092068-A1.
 PD 15-MAY-2003.
 PA (ITOH/) ITOH A.
 PA (HANA/) HANAZONO Y.
 PA (OKAD/) OKADA T.
 PA (OZAW/) OZAWA K.
 Query Match
 Best Local Similarity 25.9%; Score 156.5; DB 6; Length 412;
 RESULT 691
 ID ABG02019 standard; protein; 737 AA.
 DE Novel human diagnostic protein #2010.
 PN WO200175067-A2.
 PD 11-OCT-2001.
 PA (HYSE-) HYSEQ INC. 9.2%; Score 156; DB 4; Length 737;
 Best Local Similarity 24.3%; Pred. No. 0.00031;
 RESULT 692
 ID ADR66297 standard; protein; 358 AA.
 DE Human prostatic carcinoma derived protein SEQ ID 151 #2.
 PN WO2004076614-A2.
 PD 10-SEP-2004.
 PA (HINZ/) HINZMANN B.
 PA (DAHL/) DAHL E.
 PA (ROSE/) ROSENTHAL A.
 PA (HERM/) HERMANN K.
 PA (PILA/) PILARSKY C.
 Query Match
 Best Local Similarity 23.1%; Score 153.5; DB 8; Length 358;
 RESULT 693
 ID ADR66858 standard; protein; 358 AA.
 DE Human prostatic carcinoma derived DNA SEQ ID 151 #4.
 PN WO2004076614-A2.
 PD 10-SEP-2004.
 PA (HINZ/) HINZMANN B.
 PA (DAHL/) DAHL E.
 PA (ROSE/) ROSENTHAL A.
 PA (HERM/) HERMANN K.
 PA (PILA/) PILARSKY C.
 Query Match
 Best Local Similarity 23.1%; Score 153.5; DB 8; Length 358;
 RESULT 694
 ID AA41692 standard; protein; 373 AA.
 DE Human PRO 363 protein sequence.
 PN WO9946281-A2.
 PD 16-SEP-1999.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 2; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 695
 ID AAB33430 standard; protein; 373 AA.

DE Human PRO363 protein UNQ318 SEQ ID NO:87.
 PN WO200053758-A2.
 PD 14-SEP-2000.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 3; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 696
 ID AAB4248 standard; protein; 373 AA.
 DE Human PRO363 (UNQ318) protein sequence SEQ ID NO:59.
 PN WO200053756-A2.
 PD 14-SEP-2000.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 3; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 697
 ID AAU12365 standard; protein; 373 AA.
 DE Human PRO363 polypeptide sequence.
 PN WO200140466-A2.
 PD 07-JUN-2001.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 4; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 698
 ID AAB48146 standard; protein; 373 AA.
 DE Human A236 variant 2 polypeptide.
 PN WO200069885-A2.
 PD 23-NOV-2000.
 PA (MILL-) MILLENNIUM PHARM INC. 9.1%; Score 153.5; DB 4; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 699
 ID AAB48108 standard; protein; 373 AA.
 DE Human A236 polypeptide.
 PN WO200069885-A2.
 PD 23-NOV-2000.
 PA (MILL-) MILLENNIUM PHARM INC. 9.1%; Score 153.5; DB 4; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 700
 ID AAB65293 standard; protein; 373 AA.
 DE Human PRO363 protein sequence SEQ ID NO:503.
 PN WO200073454-A1.
 PD 07-DEC-2000.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 4; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 701
 ID AAU83656 standard; protein; 373 AA.
 DE Human PRO protein, Seq ID No 130.
 PN WO200208288-A2.
 PD 31-JAN-2002.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 5; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 702
 ID ABB84848 standard; protein; 373 AA.
 DE Human PRO363 protein sequence SEQ ID NO:64.
 PN WO200200690-A2.
 PD 03-JAN-2002.
 PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 5; Length 373;
 Query Match
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 703
 ID AAE26448 standard; protein; 373 AA.
 DE Human A236 protein.
 PN US2002055139-A1.
 PD 09-MAY-2002.
 PA (HOLT/) HOLTZMAN D A.
 PA (SHAR/) SHARP J D.
 PA (LEIB/) LEIBY K R.
 PA (BOSS/) BOSSONE S.
 PA (PANY/) PAN Y.
 PA (BARN/) BARNES T M.

PA (FRAS/) FRASER C C.
PA (WRIG/) WRIGHTON N.
PA (MYER/) MYERS P S.
PA (KING/) KINGSBURY G.
Query Match 9.1%; Score 153.5; DB 5; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 704
ID ABB95454 standard; protein; 373 AA.
DE Human angiogenesis related protein PRO363 SEQ ID NO: 64.
PN W0200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
PA (BAKE/) BAKER K P.
PA (FERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (STEP/) STEPHAN J P.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 9.1%; Score 153.5; DB 5; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 705
ID ABU59108 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003027163-A1.
PD 06-FEB-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 706
ID ABU59186 standard; protein; 373 AA.
DE Novel human secreted or transmembrane protein PRO363.
PN US2002132252-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 707
ID ABU82698 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003032023-A1.
PD 13-FEB-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 708
ID ABO17809 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003032156-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 709
ID ABU60617 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2002160384-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 710
ID ABU80803 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003036635-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 711
ID ABO25194 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003050239-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 712
ID ABO33769 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003045687-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 713
ID ABU13999 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2002103125-A1.
PD 01-AUG-2002.
PA (GETH) GENENTECH LTD.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 714
ID ABU81063 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003004311-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 715
ID ABU72200 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2002192706-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 716
ID ABU72584 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003003531-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 717
ID ABU66763 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003036180-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 718
ID ABU84880 standard; protein; 373 AA.
DE Human secreted and transmembrane polypeptide PRO363.
PN US2002177553-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 719
ID ABU59844 standard; protein; 373 AA.
DE Novel secreted and transmembrane protein PRO363.
PN US2003017563-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 720
ID ABU61078 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2002169284-A1.
PD 14-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 721
ID ABUS9333 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2003027162-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 722
ID ABO26030 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2002127576-A1.
PD 12-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 723
ID ABO25034 standard; protein; 373 AA.
DE Human secreted/transmembrane protein (PRO) #194.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 724
ID ABUS9347 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003004102-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 725
ID ABUS9039 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 726
ID ABUS9039 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 727
ID ABUS9417 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003022187-A1.
PD 30-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 728
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted or transmembrane protein PRO5723.
PN US2003027985-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 729
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted or transmembrane protein PRO5723.
PN US2003027985-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 730
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 731
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2002123463-A1.
PD 05-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 732
ID ABUS9482 standard; protein; 373 AA.
DE Human secreted and transmembrane protein PRO363.
PN US200217164-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 733
ID ABUS9482 standard; protein; 373 AA.
DE Human secreted and transmembrane polypeptide PRO363.
PN US2002197615-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 734
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003017981-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 735
ID AD445907 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 736
ID AD445907 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 737
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO363 protein.
PN US2003050448-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 738
ID AD445907 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.

PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 730
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 731
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2002123463-A1.
PD 05-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 732
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US200217164-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 733
ID ABUS9482 standard; protein; 373 AA.
DE Human secreted and transmembrane polypeptide PRO363.
PN US2002197615-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 734
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003017981-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 735
ID AD445907 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 736
ID AD445907 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 737
ID ABUS9482 standard; protein; 373 AA.
DE Human PRO363 protein.
PN US2003050448-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
RESULT 738
ID AD445907 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 739
ID ADA61611 standard; protein; 373 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 740
ID ADB19396 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 741
ID ADB27937 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 742
ID ADA86416 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 743
ID ADB15980 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087350-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 744
ID ADA38014 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US200308297-A1.
PD 09-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 745
ID ADA47766 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 746
ID ADA21700 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003054404-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 747
ID ADA10487 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003059831-A1.
PD 27-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 748

ID ADA67561 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 749
ID ADB30568 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 750
ID ADA85864 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082693-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 751
ID ADA18031 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003054987-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 752
ID ADA97076 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 753
ID ADA79380 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 754
ID ADA87519 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087345-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 755
ID ADB16721 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 756
ID ADA28139 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003054359-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 757
ID ADA91813 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082694-A1.

PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 758
ID ADB14876 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087451-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 759
ID ADA24598 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003050241-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 760
ID ADB18837 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 761
ID ADA94052 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 762
ID ADB19948 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082691-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 763
ID ADB13260 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 764
ID ABO43342 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003044945-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 765
ID ABO19649 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003050240-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 766
ID ADA12259 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003055216-A1.
PD 20-MAR-2003.

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 767
ID ADA94719 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059832-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 768
ID ADA74514 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 769
ID ADB24747 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077713-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 770
ID ADA82271 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 771
ID ADA75234 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 772
ID ADA85312 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 773
ID ADA84760 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 774
ID ADB30016 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 775
ID ADA80544 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 776
ID ADA12259 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003055216-A1.
PD 20-MAR-2003.


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Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 776
ID ADA75786 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 777
ID ADA38944 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059780-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 778
ID ADA47011 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 779
ID ADB25307 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077715-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 780
ID ADA93483 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077721-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 781
ID ADB26833 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 782
ID ADB31120 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 783
ID ABJ72420 standard; protein; 373 AA.
DE Human PRO363 protein.
PN US2003027988-A1.
PD 06-FEB-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 784
ID ADA93065 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003060407-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 785
ID ADA61048 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068797-A1.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 786
ID ADB24195 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077714-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 787
ID ADA96524 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082890-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 788
ID ADA81096 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 789
ID ADA95972 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082759-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 790
ID ADB26281 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 791
ID ADB21766 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 792
ID ABO34315 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO 363.
PN US2003044934-A1.
PD 06-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 793
ID ABO19540 standard; protein; 373 AA.
DE Novel human secreted and transmembrane polypeptide #8.
PN US2003049633-A1.
PD 13-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 794
ID ADA77545 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068797-A1.
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PD 10-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 795
ID ADB18285 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 796
ID ADA86968 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 797
ID ADA88071 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 798
ID ADA46459 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 799
ID ADB28489 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 800
ID ADB29041 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 801
ID ABO53245 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003044806-A1.
PD 06-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 802
ID ADA76993 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 803
ID ADA22626 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003040473-A1.
PD 27-FEB-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 804
ID ADA8623 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 805
ID ADA97628 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 806
ID ADB27385 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003022239-A1.
PD 30-JAN-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 807
ID ADB22318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087344-A1.
PD 08-MAY-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 808
ID ABO22615 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003017982-A1.
PD 23-JAN-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 809
ID ADA06792 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003049638-A1.
PD 13-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 810
ID ABJ72122 standard; protein; 373 AA.
DE Human membrane bound receptor/protein PRO363 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 811
ID ADA39485 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059782-A1.
PD 27-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 812
ID ADA67009 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 813
ID ADB22870 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077711-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 814
ID ADB23643 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 815
ID ADA92365 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 816
ID ADB15428 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 817
ID ADB93620 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073814-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 818
ID ADB80726 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088068-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 819
ID ADB73267 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096968-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 820
ID ADB38680 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 821
ID ADB96511 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003054403-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 822
ID ADB78349 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092889-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 823
ID ADB38128 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 824
ID ADB66600 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 825
ID ADB84997 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003073817-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 826
ID ADB89680 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 827
ID ADB90412 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 828
ID ADB39513 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 829
ID ADB78103 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092886-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 830
ID ADB73565 standard; protein; 373 AA.
DE Human PRO polypeptide #8.
PN US2003045462-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 831
ID ADB87169 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003088067-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
RESULT 832
ID ADB78349 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092889-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;

RESULT 832
ID ADB84751 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003092890-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 833
ID ADB47136 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082887-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 834
ID ADB83866 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003069397-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 835
ID ADB86743 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082697-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 836
ID ADB73021 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092887-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 837
ID ADB76281 standard; protein; 373 AA.
DE Human PRO polypeptide #8.
PN US2003083248-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 838
ID ADB77348 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082896-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 839
ID ADB34505 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077717-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 840
ID ADB35609 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 841
ID ADB35609 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;

ID	ADB33953 standard; protein; 373 AA.
DE	Human PRO polypeptide SEQ ID NO 388.
FN	US2003077716-A1.
PD	24-APR-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 842	
ID	ADB35057 standard; protein; 373 AA.
DE	Human PRO polypeptide SEQ ID NO 388.
FN	US2003077718-A1.
PD	24-APR-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 843	
ID	ADB36161 standard; protein; 373 AA.
DE	Human PRO polypeptide SEQ ID NO 388.
FN	US2003077720-A1.
PD	24-APR-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 844	
ID	ADB46556 standard; protein; 373 AA.
DE	Novel human secreted and transmembrane protein PRO363.
FN	US2003082692-A1.
PD	01-MAY-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 845	
ID	ADC43707 standard; protein; 373 AA.
DE	Human secreted/transmembrane protein, PRO363.
FN	US2003054586-A1.
PD	20-MAR-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 846	
ID	ADC57983 standard; protein; 373 AA.
DE	Human PRO polypeptide #140.
FN	US2003027754-A1.
PD	06-FEB-2003.
PA	
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 847	
ID	ADC55347 standard; protein; 373 AA.
DE	Human PRO polypeptide #140.
FN	US2003045463-A1.
PD	06-MAR-2003.
PA	
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 848	
ID	ADC12214 standard; protein; 373 AA.
DE	Human secreted/transmembrane protein PRO363.
FN	US2003049681-A1.
PD	13-MAR-2003.
PA	
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 849	
ID	ADC61467 standard; protein; 373 AA.
DE	Human secreted/transmembrane protein, PRO363.
FN	US2003049684-A1.
PD	13-MAR-2003.
PA	(GETH) GENENTECH INC.
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;
RESULT 850	
ID	ADC63431 standard; protein; 373 AA.
DE	Human secreted/transmembrane protein, PRO363.
FN	US2003054405-A1.
PD	20-MAR-2003.
PA	
Query Match	9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity	27.0%; Pred. No. 0.0002;

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 851
ID ADC66531 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003060406-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 852
ID ADC56636 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003064375-A1.
PD 03-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 853
ID ADC68655 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003064407-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 854
ID ADC62715 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003068648-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 855
ID ADC67780 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003069178-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 856
ID ADC11681 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003069403-A1.
PD 10-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 857
ID ADC41100 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003072745-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 858
ID ADC67155 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003073331-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 859
ID ADC62091 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003073624-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 860
ID ADC36859 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US200308065-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 861
ID ADC41724 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003104998-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 862
ID ADC21849 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096969-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 863
ID ADC50429 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092106-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 864
ID ADC71976 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092107-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 865
ID ADC59955 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092105-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 866
ID ADC49880 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088064-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 867
ID ADC49079 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088070-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 868
ID ADC49596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 869
ID ADC49596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;

ID ADC47457 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088072-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 870
ID ADC52962 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087365-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 871
ID ADC57316 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 872
ID ADC60507 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087367-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 873
ID ADC50982 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 874
ID ADC65509 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 875
ID ADC54607 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 876
ID ADC53568 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 877
ID ADC59091 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 878
ID ADC55969 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 879
ID ADC58539 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 880
ID ADC14803 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082546-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 881
ID ADC47202 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105288-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 882
ID ADD08335 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003068623-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 883
ID ADD03213 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092104-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 884
ID ADC90205 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 885
ID ADC82160 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003083461-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 886
ID ADC69624 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194770-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 887
ID ADC48513 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 888
ID ADC47961 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 889
ID ADD07802 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2002193299-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 890
ID ADC78077 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096972-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 891
ID ADD04617 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087354-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 892
ID ADC82693 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003059833-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 893
ID ADD06312 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073816-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 894
ID ADC80573 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 895
ID ADD11080 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 896
ID ADD10353 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;

RESULT 897
ID ADC47961 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 898
ID ADD08873 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073090-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 899
ID ADC77831 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088066-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 900
ID ADC80021 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 901
ID ADD07122 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2002193300-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 902
ID ADD11313 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 903
ID ADD09490 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 904
ID ADC83369 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003059783-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 905
ID ADD50794 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105291-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 906
ID ADD41203 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.

PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 907
ID ADD52342 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 908
ID ADD51040 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105290-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 909
ID ADD53082 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194792-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 910
ID ADD53634 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003203437-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 911
ID ADD5476 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003077593-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 912
ID ADD37106 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 913
ID ADD56434 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003077594-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 914
ID ADD51790 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 915
ID ADD02589 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;

Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 916
ID ADD50521 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096971-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 917
ID ADD02023 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 918
ID ADD54205 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 919
ID ADD54872 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2002132253-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 920
ID ADD50275 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096970-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 921
ID ADD51286 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105289-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 922
ID ADD49093 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003096744-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 923
ID ADD92522 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 924
ID ADD91418 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 925
ID ADE04032 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199057-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 926
ID ADE31891 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003086647-A1.
PD 10-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 927
ID ADE27026 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087304-A1.
PD 08-MAY-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 928
ID ADE33229 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 929
ID ADE22261 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 930
ID ADD79485 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 931
ID ADE35147 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203434-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 932
ID ADE16261 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203435-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 933
ID ADD72876 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203436-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 934
ID ADE42021 standard; protein; 373 AA.

DE Human PRO polypeptide #194.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 935
ID ADE17838 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 936
ID ADD91970 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 937
ID ADE33433 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194767-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 938
ID ADE33985 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 939
ID ADD80037 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207417-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 940
ID ADD93074 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194768-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 941
ID ADD72234 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194781-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 942
ID ADE19494 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 943
ID ADE18942 standard; protein; 373 AA.
DE Human PRO polypeptide #194.

PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 944
ID ADE43138 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 945
ID ADD95927 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 946
ID ADE22813 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199064-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 947
ID ADD78931 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203429-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 948
ID ADE26493 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087305-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 949
ID ADE32881 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 950
ID ADE42573 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 951
ID ADE16885 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203433-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 952
ID ADD80589 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207418-A1.
PD 06-NOV-2003.

PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 953
ID ADD89617 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 954
ID ADE40901 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 955
ID ADE04700 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 956
ID ADE92829 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 957
ID ADF46899 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195333-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 958
ID ADF67430 standard; protein; 373 AA.
DE Human PRO363 amino acid sequence SEQ ID NO:503.
PN US2002198148-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 959
ID ADG21538 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207355-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 960
ID ADG23179 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 961
ID ADF97514 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 962
ID ADG80578 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 963
ID ADG52656 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003216561-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 964
ID ADG59976 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003206915-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 965
ID ADG80026 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 966
ID ADH55318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 967
ID ADH55870 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 968
ID ADI35684 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003050457-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 969
ID ADI60736 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003077000-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 970
ID ADI64089 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 971
ID ADI65038 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 972
ID ADI63537 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 973
ID ADH81951 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 974
ID ADI00177 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003049682-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 975
ID ADH81399 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 976
ID ADM82568 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 977
ID ADN15967 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 978
ID ADN16596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 979
ID ADN15415 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 980
ID ADN14863 standard; protein; 373 AA.

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DE Novel human secreted and transmembrane protein PRO363.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 981
ID ADC48833 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092888-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 982
ID ADC81125 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 983
ID ADE21004 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100735-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 984
ID ADE05848 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100728-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 985
ID ADD76573 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 986
ID ADD75077 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100712-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 987
ID ADD75823 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100717-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 988
ID ADD85055 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100722-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 989
ID ADD86881 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.

PN US2003100738-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 990
ID ADE20758 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100734-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 991
ID ADE39055 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096362-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 992
ID ADD87937 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 993
ID ADD86341 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 994
ID ADE05602 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100727-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 995
ID ADD73587 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100711-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 996
ID ADE75789 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 997
ID ADE48393 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003104536-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 998
ID ADD78427 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100737-A1.

PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 999
ID ADE41314 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1000
ID ADE23365 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1001
ID ADE21250 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100736-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1002
ID ADD77365 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100732-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1003
ID ADE20512 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100733-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1004
ID ADD75577 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100084-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1005
ID ADD74093 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100708-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1006
ID ADD74339 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100709-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1007
ID ADD76069 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100718-A1.
PD 29-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1008
ID ADD85561 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100721-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1009
ID ADE23917 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1010
ID ADE24560 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1011
ID ADD87385 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1012
ID ADE05110 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100726-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1013
ID ADD75323 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100714-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1014
ID ADD76867 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100715-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1015
ID ADD86635 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100719-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1016
ID ADE89251 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1017
ID ADD78103 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100731-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1018
ID ADE18390 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194794-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1019
ID ADE86699 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199054-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1020
ID ADE89494 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003130181-A1.
PD 10-JUL-2003.
PA (ASHK/) ASHKENAZI A J.
PA (BAKE/) BAKER K P.
PA (BOTS/) BOTSTEIN D.
PA (DESN/) DESNOYERS L.
PA (EATO/) EATON D L.
PA (FERR/) FERRARA N.
PA (FILV/) FILVAROFF E.
PA (FONG/) FONG S.
PA (GAOW/) GAO W.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GIRM/) GIRMALDI J C.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (KLJA/) KLJAVIN I J.
PA (KUOS/) KUO S S.
PA (NAPI/) NAPIER M A.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (ROYM/) ROY M A.
PA (SHEL/) SHELTON D L.
PA (STEW/) STEWART T A.
PA (TUMA/) TUMAS D.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1021
ID ADD77611 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100729-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1022
ID ADD77857 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100730-A1.
PD 29-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1023
ID ADD85315 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100725-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1024
ID ADD73947 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100710-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1025
ID ADD74585 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100713-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1026
ID ADD77113 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100716-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1027
ID ADD85807 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100720-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1028
ID ADE05356 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100723-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1029
ID ADD74831 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100724-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1030
ID ADF61134 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195345-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1031
ID ADF39826 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003198994-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

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Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1032
ID ADF45622 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195148-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1033
ID ADE94719 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1034
ID ADE91130 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1035
ID ADF35629 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003194760-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1036
ID ADE95271 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1037
ID ADE93381 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1038
ID ADF24018 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003204055-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1039
ID ADF40450 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1040
ID ADF23394 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203402-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1041
ID ADF33377 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194780-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1042
ID ADF34962 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1043
ID ADF26844 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199436-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1044
ID ADF27480 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199437-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1045
ID ADE92277 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1046
ID ADE90578 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1047
ID ADF41074 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199435-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1048
ID ADF32753 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003211091-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1049
ID ADF25119 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003211092-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1050
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ID ADF26220 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199674-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1051
ID ADF34009 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194410-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1052
ID ADF46246 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195344-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1053
ID ADE91725 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1054
ID ADG11879 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003228655-A1.
PD 11-DEC-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1055
ID ADG05643 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096959-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1056
ID ADG27197 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096962-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1057
ID ADG02304 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1058
ID ADG22090 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1059
ID ADG20160 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1060
ID ADF98066 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1061
ID ADG24283 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1062
ID ADF98637 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1063
ID ADG03468 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1064
ID ADF99189 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1065
ID ADG16774 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1066
ID ADG05233 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1067
ID ADG19500 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1068
ID ADG11260 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
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PN US2003096967-A1.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1069
ID ADG13337 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1070
ID ADG08394 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1071
ID ADG15564 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1072
ID ADG12039 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096963-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1073
ID ADF96962 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1074
ID ADG06147 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1075
ID ADG23731 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1076
ID ADG04020 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1077
ID ADG24921 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207427-A1.

PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1078
ID ADF94596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096964-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1079
ID ADG07218 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1080
ID ADG07770 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1081
ID ADG0692 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096966-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1082
ID ADG55265 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1083
ID ADG60929 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1084
ID ADG2033 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1085
ID ADG82234 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1086
ID ADG57473 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207362-A1.
PD 06-NOV-2003.

Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1096				
ID ADG81682 standard; protein; 373 AA.				
DE Human PRO polypeptide #194.				
FN US2003207805-A1.				
PD 06-NOV-2003.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1097				
ID ADH19749 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein PRO363.				
FN US2003228656-A1.				
PD 11-DEC-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1098				
ID ADH30644 standard; protein; 373 AA.				
DE Human PRO polypeptide #194.				
FN US2003077723-A1.				
PD 24-APR-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1099				
ID ADH12011 standard; protein; 373 AA.				
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207419-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1100				
ID ADG49608 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003215905-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1101				
ID ADG51480 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003215908-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1102				
ID ADG52433 standard; protein; 373 AA.				
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207414-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1103				
ID ADG54161 standard; protein; 373 AA.				
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207416-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1104				
ID ADG48984 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1105				
ID ADG48984 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1106				
ID ADG48984 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1107				
ID ADG48984 standard; protein; 373 AA.				
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.	9.1%;	Score 153.5;	DB 8;	Length 373;
Query Match	27.0%;	Pred. No. 0.0002;		
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1108				

RESULT 1105
ID ADG81130 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194793-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1106
ID ADG56369 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207366-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1107
ID ADH12635 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207378-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1108
ID ADG48360 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003216560-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1109
ID ADH21242 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003224358-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1110
ID ADG61481 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207429-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1111
ID ADH20282 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003219856-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1112
ID ADH28568 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003022331-A1.
PD 30-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1113
ID ADG54713 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207367-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1114
ID ADG59753 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein PRO363.
PN US2003207369-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1115
ID ADG50856 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004005312-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1116
ID ADH43497 standard; protein; 373 AA.
DE Human PRO polypeptide #32.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1117
ID ADG58800 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004005657-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1118
ID ADG34126 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004006206-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1119
ID ADG62256 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004006219-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1120
ID ADH1177 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1121
ID ADI33596 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096960-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1122
ID ADH69690 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2004019183-A1.
PD 29-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1123
ID ADH25281 standard; protein; 373 AA.
DE Human neurotrophin homologue related protein sequence SEQ ID NO:59.

PD EPI386931-A1.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1124
ID ADG09920 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004009548-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1125
ID ADI15391 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207382-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1126
ID ADG09268 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004009547-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1127
ID ADI14723 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207383-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1128
ID ADI29851 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096961-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1129
ID ADI18318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207349-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1130
ID ADM27248 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US200404179-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1131
ID ADJ63599 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004039164-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1132
ID ADJ77494 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004038336-A1.

PD 26-FEB-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1133
ID ADK82842 standard; protein; 373 AA.
DE Human PRO polypeptide #32.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1134
ID ADK66606 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2004044180-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1135
ID ADJ65616 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004038335-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1136
ID ADM27752 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004048333-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1137
ID ADM17058 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004048332-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1138
ID ADJ06892 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004063921-A1.
PD 01-APR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1139
ID ADM42476 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004058424-A1.
PD 25-MAR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1140
ID ADN05373 standard; protein; 373 AA.
DE Antipsoriatic protein sequence #858.
PN WO2004028479-A2.
PD 08-APR-2004.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1141
ID ADM28338 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004077064-A1.
PD 22-APR-2004.

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1142
ID ADO36720 standard; protein; 373 AA.
DE Human UKW polypeptide, SEQ ID NO:2.
PN EP1416279-A1.
PD 06-MAY-2004.
PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1143
ID ADI95820 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077659-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1144
ID ADI96372 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207354-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1145
ID ADA54925 standard; protein; 512 AA.
DE Human protein, SEQ ID 2493.
PN EP1293589-A2.
PD 19-MAR-2003.
PA (HELI-) HELIX RES INST.
Query Match 9.1%; Score 153.5; DB 6; Length 512;
Best Local Similarity 25.2%; Pred. No. 0.00031;
RESULT 1146
ID AAY53666 standard; protein; 4412 AA.
DE Sequence gi/1017427/emb/CAAG2189 from an alignment with protein 608.
PN WO9960164-A1.
PD 25-NOV-1999.
PA (QUAR-) QUARK BIOTECH INC.
Query Match 9.1%; Score 153.5; DB 3; Length 4412;
Best Local Similarity 22.0%; Pred. No. 0.0058;
RESULT 1147
ID ABP60991 standard; protein; 5635 AA.
DE Novel human protein. SEQ ID 78.
PN WO200250105-A1.
PD 27-JUN-2002.
PA (SMIK) SMITHKLINE BEECHAM CORP.
PA (SMIK) SMITHKLINE BEECHAM PLC.
PA (GLAX) GLAXO GROUP LTD.
Query Match 9.1%; Score 153.5; DB 5; Length 5635;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1148
ID ADJ70089 standard; protein; 5636 AA.
DE Human heat mitochondrial protein as a therapeutic target SeqID1895.
PN WO2003087768-A2.
PD 23-OCT-2003.
PA (MITO-) MITOKOR.
PA (BUCK-) BUCK INST AGE RES.
Query Match 9.1%; Score 153.5; DB 7; Length 5636;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1149
ID ADJ83137 standard; protein; 5636 AA.
DE Human hemicentin protein - SEQ ID 128.
PN US2003170630-A1.
PD 11-SEP-2003.
PA (ALSO/) ALSOBROOK J P.
PA (TCHV/) TCHERNEV V T.
PA (LIUX/) LIU X.
PA (SPYT/) SPYTEK K A.
PA (ZERR/) ZERRHUSEN B D.

PA (PATT/) PATTURAJAN M.
PA (LEPL/) LEPLEY D M.
PA (BURG/) BURGESS C E.
PA (SHIM/) SHIMKETS R A.
PA (GROS/) GROSSE W M.
PA (SZEK/) SZEKERES E S.
PA (VERN/) VERNET C A M.
PA (LILL/) LI L.
PA (CASM/) CASMAN S J.
PA (BOLD/) BOLDOG F L.
PA (GORM/) GORMAN L.
PA (GANG/) GANGOLLI E A.
PA (FERN/) FERNANDES E R.
PA (RIEG/) RIEGER D K.
PA (EDIN/) EDINGER S R.
PA (GUNT/) GUNTHER E.
PA (MILL/) MILLET I.
PA (SCIO/) SCIORE P.
PA (ELLE/) ELLERMAN K.
PA (MACD/) MACDOUGALL J R.
PA (SMIT/) SMITHSON G.
Query Match 9.1%; Score 153.5; DB 7; Length 5636;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1150
ID ADK60205 standard; protein; 5636 AA.
DE Angiogenesis differentially expressed protein GS-P29.
PN FR2836687-A1.
PD 05-SEP-2003.
PA (GENE-) GENE SIGNAL.
PA (ALMA/) AL MAHMOOD S.
Query Match 9.1%; Score 153.5; DB 8; Length 5636;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1151
ID ADK60506 standard; protein; 5636 AA.
DE Angiogenesis differentially expressed protein GS-P29.
PN FR2836686-A1.
PD 05-SEP-2003.
PA (GENE-) GENE SIGNAL.
PA (ALMA/) AL MAHMOOD S.
Query Match 9.1%; Score 153.5; DB 8; Length 5636;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1152
ID ADP73129 standard; protein; 5636 AA.
DE Angiogenesis inhibitor human protein sequence, GS-P29.
PN FR2843753-A1.
PD 27-FEB-2004.
PA (GENE/) GENE S.
PA (ALMS/) AL M S.
Query Match 9.1%; Score 153.5; DB 8; Length 5636;
Best Local Similarity 25.2%; Pred. No. 0.0081;
RESULT 1153
ID ADA50158 standard; protein; 393 AA.
DE Human CAR/mouse SCF mature fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 9.1%; Score 153; DB 6; Length 393;
Best Local Similarity 24.3%; Pred. No. 0.00024;
RESULT 1154
ID ABG74786 standard; protein; 31267 AA.
DE Human RGS11 protein.
PN WO2002103355-A1.
PD 27-DEC-2002.
PA (TAKE) TAKEDA CHEM IND LTD.
Query Match 9.1%; Score 153; DB 6; Length 31267;
Best Local Similarity 28.2%; Pred. No. 0.091;
RESULT 1155
ID ADA50159 standard; protein; 474 AA.
DE Human CAR/mouse anti-CD34 antibody mature fusion protein.
PN US2003092068-A1.

PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 9.0%; Score 152; DB 6; Length 474;
Best Local Similarity 23.2%; Pred. No. 0.00038;
RESULT 1156
ID AAB48145 standard; protein; 373 AA.
DE Human A236 variant 1 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 9.0%; Score 151.5; DB 4; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0003;
RESULT 1157
ID AAU17996 standard; protein; 301 AA.
DE Human immunoglobulin polypeptide SEQ ID NO 141.
PN WO20015315-A2.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.9%; Score 151; DB 4; Length 301;
Best Local Similarity 25.8%; Pred. No. 0.00025;
RESULT 1158
ID ABB10232 standard; protein; 301 AA.
DE Human cDNA SEQ ID NO: 540.
PN WO200154474-A2.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.9%; Score 151; DB 4; Length 301;
Best Local Similarity 25.8%; Pred. No. 0.00025;
RESULT 1159
ID ABP66819 standard; protein; 301 AA.
DE Human polypeptide SEQ ID NO 540.
PN US2002090672-A1.
PD 11-JUL-2002.
PA (ROSE/) ROSEN C A.
PA (RUBE/) RUBEN S M.
PA (BARA/) BARASH S C.
Query Match 8.9%; Score 151; DB 5; Length 301;
Best Local Similarity 25.8%; Pred. No. 0.00025;
RESULT 1160
ID ADB31620 standard; protein; 301 AA.
DE Human novel protein SEQ ID NO 141.
PN US2003077606-A1.
PD 24-APR-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.9%; Score 151; DB 7; Length 301;
Best Local Similarity 25.8%; Pred. No. 0.00025;
RESULT 1161
ID ADR41522 standard; protein; 318 AA.
DE Human CD-like molecule HKAC103, SEQ ID NO:321.
PN WO200226930-A2.
PD 04-APR-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.9%; Score 151; DB 5; Length 318;
Best Local Similarity 25.8%; Pred. No. 0.00027;
RESULT 1162
ID ADA50157 standard; protein; 393 AA.
DE Human CAR/SCF mature fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 8.9%; Score 151; DB 6; Length 393;
Best Local Similarity 25.4%; Pred. No. 0.00036;
RESULT 1163
ID ABG02117 standard; protein; 434 AA.
DE Novel human diagnostic protein #2108.
PN WO200175067-A2.
PD 11-OCT-2001.

PA (HYSE-) HYSEQ INC.
Query Match 8.9%; Score 151; DB 4; Length 434;
Best Local Similarity 24.3%; Pred. No. 0.00041;
RESULT 1164
ID ABU62399 standard; protein; 466 AA.
DE Chimeric CAR/Hg/Pro-A gene product.
PN US6524572-B1.
PD 25-FEB-2003.
PA (RAIN-) RAINBOW THERAPEUTIC CO.
Query Match 8.9%; Score 151; DB 6; Length 466;
Best Local Similarity 25.4%; Pred. No. 0.00045;
RESULT 1165
ID AAB48147 standard; protein; 373 AA.
DE Human A236 variant 3 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 8.9%; Score 150.5; DB 4; Length 373;
Best Local Similarity 26.6%; Pred. No. 0.00037;
RESULT 1166
ID ABG31321 standard; protein; 2652 AA.
DE Predicted human adican-2 protein.
PN WO200246364-A2.
PD 13-JUN-2002.
PA (QUAR-) QUARK BIOTECH INC.
Query Match 8.7%; Score 147.5; DB 5; Length 2652;
Best Local Similarity 21.8%; Pred. No. 0.0095;
RESULT 1167
ID ADU02250 standard; protein; 2652 AA.
DE Human OCP protein #6.
PN US2004053301-A1.
PD 18-MAR-2004.
PA (QUAR-) QUARK BIOTECH INC.
Query Match 8.7%; Score 147.5; DB 8; Length 2652;
Best Local Similarity 21.8%; Pred. No. 0.0095;
RESULT 1168
ID ABP70049 standard; protein; 2845 AA.
DE Human NOV1a.
PN WO200272771-A2.
PD 19-SEP-2002.
PA (CURA-) CURAGEN CORP.
Query Match 8.7%; Score 147.5; DB 5; Length 2845;
Best Local Similarity 21.8%; Pred. No. 0.01;
RESULT 1169
ID ABG22401 standard; protein; 361 AA.
DE Novel human diagnostic protein #22392.
PN WO200175067-A2.
PD 11-OCT-2001.
PA (HYSE-) HYSEQ INC.
Query Match 8.7%; Score 146.5; DB 4; Length 361;
Best Local Similarity 24.5%; Pred. No. 0.00078;
RESULT 1170
ID AAB85862 standard; protein; 373 AA.
DE Murine adipocytes-derived protein.
PN WO200166720-A1.
PD 13-SEP-2001.
PA (KITA/) KITAMURA T.
PA (TSUR/) TSURUGA H.
Query Match 8.7%; Score 146.5; DB 4; Length 373;
Best Local Similarity 25.9%; Pred. No. 0.00081;
RESULT 1171
ID AAB48126 standard; protein; 373 AA.
DE Mouse A236 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 8.6%; Score 145.5; DB 4; Length 373;
Best Local Similarity 25.9%; Pred. No. 0.00099;
RESULT 1172
ID AAE26449 standard; protein; 373 AA.
DE Mouse A236 protein.
PN US2002055139-A1.
PD 09-MAY-2002.

PA (HOLT/) HOLTZMAN D A.
PA (SHAR/) SHARP J D.
PA (LEIB/) LEIBY K R.
PA (BOSS/) BOSSONE S.
PA (PANY/) PAN Y.
PA (BAEN/) BARNES T M.
PA (FRAS/) FRASER C C.
PA (WRIG/) WRIGHTON N.
PA (MYER/) MYERS P S.
PA (KING/) KINGSBURY G.
Query Match 8.6%; Score 145.5; DB 5; Length 373;
Best Local Similarity 25.9%; Pred. No. 0.00099;
RESULT 1173
ID ABB66424 standard; protein; 2016 AA.
DE Drosophila melanogaster polypeptide SEQ ID NO 26064.
PN WO200171042-A2.
PD 27-SEP-2001.
PA (PEKE) PE CORP NY.
Query Match 8.6%; Score 145; DB 4; Length 2016;
Best Local Similarity 25.6%; Pred. No. 0.011;
RESULT 1174
ID AAB48149 standard; protein; 373 AA.
DE Mouse A236 variant 2 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 26.1%; Pred. No. 0.0012;
RESULT 1175
ID AAB48150 standard; protein; 373 AA.
DE Mouse A236 variant 3 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 25.6%; Pred. No. 0.0012;
RESULT 1176
ID AAB48148 standard; protein; 373 AA.
DE Mouse A236 variant 1 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 26.1%; Pred. No. 0.0012;
RESULT 1177
ID ADG63213 standard; protein; 367 AA.
DE Human neurotrophin protein +69bp isoform.
PN WO2003002765-A2.
PD 09-JAN-2003.
PA (IMCR) IMPERIAL CANCER RES TECHNOLOGY LTD.
Query Match 8.5%; Score 144; DB 7; Length 367;
Best Local Similarity 26.6%; Pred. No. 0.0013;
RESULT 1178
ID ABB63044 standard; protein; 467 AA.
DE Drosophila melanogaster polypeptide SEQ ID NO 15924.
PN WO200171042-A2.
PD 27-SEP-2001.
PA (PEKE) PE CORP NY.
Query Match 8.5%; Score 143.5; DB 4; Length 467;
Best Local Similarity 22.1%; Pred. No. 0.002;
RESULT 1179
ID ADE37401 standard; protein; 396 AA.
DE Murine nectin-like protein 1.
PN WO2003064992-A2.
PD 07-AUG-2003.
PA (PLAN-) PLANET BIOTECHNOLOGY INC.
PA (LARR/) LARRICK J W.
PA (WYCO/) WYCOFF K L.
Query Match 8.4%; Score 142.5; DB 7; Length 396;
Best Local Similarity 19.6%; Pred. No. 0.0019;
RESULT 1180
ID AAY69287 standard; protein; 398 AA.
DE Amino acid sequence of long extracellular form of murine B7-1 (CD80).

PN WO200008057-A2.
PD 17-FEB-2000.
PA (IMVX) IMMUNEX CORP.
Query Match 8.4%; Score 142.5; DB 3; Length 398;
Best Local Similarity 19.6%; Pred. No. 0.0019;
RESULT 1181
ID AAE00868 standard; protein; 404 AA.
DE Mouse brain immunoglobulin superfamily receptor (Bigr) protein.
PN WO200129083-A1.
PD 26-APR-2001.
PA (TEXA-) TEXAS BIOTECHNOLOGY CORP.
Query Match 8.4%; Score 142.5; DB 4; Length 404;
Best Local Similarity 19.6%; Pred. No. 0.002;
RESULT 1182
ID ABB68257 standard; protein; 1395 AA.
DE Drosophila melanogaster polypeptide SEQ ID NO 31563.
PN WO200171042-A2.
PD 27-SEP-2001.
PA (PEKE) PE CORP NY.
Query Match 8.4%; Score 142.5; DB 4; Length 1395;
Best Local Similarity 21.1%; Pred. No. 0.011;
RESULT 1183
ID ADE08000 standard; protein; 376 AA.
DE Novel protein (useful for identifying genetic disorders) #155.
PN WO2003054152-A2.
PD 03-JUL-2003.
PA (HYSE-) HYSEQ INC.
Query Match 8.4%; Score 141.5; DB 7; Length 376;
Best Local Similarity 25.2%; Pred. No. 0.0022;
RESULT 1184
ID AAY13563 standard; protein; 1395 AA.
DE Drosophila Robo 1 polypeptide.
PN WO9925833-A1.
PD 27-MAY-1999.
PA (REGC) UNIV CALIFORNIA.
Query Match 8.4%; Score 141.5; DB 2; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;
RESULT 1185
ID AAY08401 standard; protein; 1395 AA.
DE Drosophila sp. Robo1 protein.
PN WO9920764-A1.
PD 29-APR-1999.
PA (REGC) UNIV CALIFORNIA.
Query Match 8.4%; Score 141.5; DB 2; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;
RESULT 1186
ID ADE85335 standard; protein; 1395 AA.
DE Fruitfly nerve cell growth modulator SLIT-1-associated sequence #1.
PN US2003170727-A1.
PD 11-SEP-2003.
PA (GOOD/) GOODMAN C S.
PA (KIDD/) KIDD T.
PA (BROS/) BROSE K.
PA (TESS/) TESSIER-LAVIGNE M.
Query Match 8.4%; Score 141.5; DB 7; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;
RESULT 1187
ID ABO84469 standard; protein; 3475 AA.
DE Human cancer-associated protein HP13-036.1.
PN WO2004074320-A2.
PD 02-SEP-2004.
PA (SAGR-) SAGRES DISCOVERY INC.
Query Match 8.4%; Score 141.5; DB 8; Length 3475;
Best Local Similarity 27.5%; Pred. No. 0.045;
RESULT 1188
ID ABB71502 standard; protein; 1052 AA.
DE Drosophila melanogaster polypeptide SEQ ID NO 41298.
PN WO200171042-A2.
PD 27-SEP-2001.
PA (PEKE) PE CORP NY.
Query Match 8.4%; Score 141; DB 4; Length 1052;
Best Local Similarity 25.1%; Pred. No. 0.0098;
RESULT 1189

ID AAW69698 standard; protein; 352 AA.
DE Mouse coxsackievirus and Ad2 and Ad5 receptor MCAR protein.
FN WO9833819-A1.
PD 06-AUG-1998.
PA (UUNY) UNIV NEW YORK STATE.
Query Match 8.3%; Score 140.5; DB 2; Length 352;
Best Local Similarity 25.4%; Pred. No. 0.0024;
RESULT 1190
ID AAW57213 standard; protein; 376 AA.
DE Mouse coxsackievirus and adenovirus receptor.
FN WO9811221-A2.
PD 19-MAR-1998.
PA (DAND) DANA FARBER CANCER INST INC.
Query Match 8.3%; Score 140.5; DB 2; Length 376;
Best Local Similarity 25.4%; Pred. No. 0.0027;
RESULT 1191
ID ADP56685 standard; protein; 265 AA.
DE Human junction adhesion molecule 3 splice variant 2 (huJAM3sv2) protein.
FN WO2004053058-A2.
PD 24-JUN-2004.
PA (ELIL) LILLY & CO ELI.
Query Match 8.3%; Score 139.5; DB 8; Length 265;
Best Local Similarity 24.0%; Pred. No. 0.002;
RESULT 1192
ID AAY96294 standard; protein; 310 AA.
DE Human IGFAM-6 immunoglobulin.
FN WO200029583-A2.
PD 25-MAY-2000.
PA (INCY-) INCYTE PHARM INC.
Query Match 8.3%; Score 139.5; DB 3; Length 310;
Best Local Similarity 22.0%; Pred. No. 0.0025;
RESULT 1193
ID ADP56683 standard; protein; 310 AA.
DE Human junction adhesion molecule 3 (huJAM3) full-length protein.
FN WO2004053058-A2.
PD 24-JUN-2004.
PA (ELIL) LILLY & CO ELI.
Query Match 8.3%; Score 139.5; DB 8; Length 310;
Best Local Similarity 22.0%; Pred. No. 0.0025;
RESULT 1194
ID AAY96735 standard; protein; 310 AA.
DE PRO1868, an A33 antigen homologue.
FN WO200036102-A2.
PD 22-JUN-2000.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1195
ID AAB33457 standard; protein; 310 AA.
DE Human PRO1868 protein UNQ859 SEQ ID NO:193.
FN WO200053758-A2.
PD 14-SEP-2000.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1196
ID AAB27276 standard; protein; 310 AA.
DE Human confluency regulated adhesion molecule 1 #2.
FN WO200053749-A2.
PD 14-SEP-2000.
PA (RMFD-) RMF DICTAGENE SA.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1197
ID AAB80272 standard; protein; 310 AA.
DE Human PRO1868 protein.
FN WO200104311-A1.
PD 18-JAN-2001.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1198
ID AAW93905 standard; protein; 310 AA.

DE Human polypeptide, SEQ ID NO: 4051.
FN EPI130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1199
ID AAM93323 standard; protein; 310 AA.
DE Human polypeptide, SEQ ID NO: 2845.
FN EPI130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1200
ID AAU12440 standard; protein; 310 AA.
DE Human PRO1868 polypeptide sequence.
FN WO200140466-A2.
PD 07-JUN-2001.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1201
ID AAB80383 standard; protein; 310 AA.
DE Secreted protein encoded by gene #13.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1202
ID AAB80408 standard; protein; 310 AA.
DE Secreted protein encoded by gene #38.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1203
ID AAB80409 standard; protein; 310 AA.
DE Secreted protein encoded by gene #39.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1204
ID ABG92709 standard; protein; 310 AA.
DE Human secreted protein PRO1868.
FN US2002098506-A1.
PD 25-JUL-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1205
ID ABG91361 standard; protein; 310 AA.
DE Novel human secreted protein #7.
FN US2002098505-A1.
PD 25-JUL-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1206
ID ABB84947 standard; protein; 310 AA.
DE Human PRO1868 protein sequence SEQ ID NO:262.
FN WO20020690-A2.
PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1207
ID ABG65297 standard; protein; 310 AA.
DE Human albumin fusion protein #1972.

PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1208
ID ABG65296 standard; protein; 310 AA.
DE Human albumin fusion protein #1971.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1209
ID ABG65298 standard; protein; 310 AA.
DE Human albumin fusion protein #1973.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1210
ID ABG31401 standard; protein; 310 AA.
DE Human PRO1868 polypeptide.
PN US2002098507-A1.
PD 25-JUL-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1211
ID ABB95553 standard; protein; 310 AA.
DE Human angiogenesis related protein PRO1868 SEQ ID NO: 262.
PN WO200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1212
ID ABB95553 standard; protein; 310 AA.
DE Human angiogenesis related protein PRO1868 SEQ ID NO: 262.
PN WO200208284-A2.
PD 31-JAN-2002.
PA (BAKE/) BAKER K P.
PA (FERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (STEP/) STEPHAN J F.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1213
ID ABU71650 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2002146709-A1.
PD 10-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1213
ID ABU72377 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2002182618-A1.
PD 05-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1214
ID ABUS0867 standard; protein; 310 AA.
DE Human secreted and transmembrane polypeptide PRO1868.
PN US2002192868-A1.
PD 19-DEC-2002.

PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1215
ID ABO17884 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003032156-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1216
ID ABU71505 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2002192659-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1217
ID ADA57610 standard; protein; 310 AA.
DE Human secreted protein #592.
PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1218
ID ADA57611 standard; protein; 310 AA.
DE Human secreted protein #592.
PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1219
ID ADA57309 standard; protein; 310 AA.
DE Human secreted protein #592.
PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1220
ID ABP71277 standard; protein; 310 AA.
DE Human junctional adhesion molecule 3 (JAM3).
PN WO2003006673-A2.
PD 23-JAN-2003.
PA (TEXA-) TEXAS BIOTECHNOLOGY CORP.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1221
ID ASU81138 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003004311-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1222
ID ABU71951 standard; protein; 310 AA.
DE Human secreted/transmembrane protein PRO1868.
PN US2003003530-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1223
ID ABO01834 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2002197671-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1224
ID ABU66838 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003036180-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1225
ID ABU54407 standard; protein; 310 AA.
DE Human secreted/transmembrane protein PRO1868.
PN US2002132240-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1226
ID AB047422 standard; protein; 310 AA.
DE Human secreted/transmembrane polypeptide PRO1868.
PN US2003044839-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1227
ID ABG73314 standard; protein; 310 AA.
DE Human PRO1868 polypeptide.
PN US2002164646-A1.
PD 07-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1228
ID ABU59919 standard; protein; 310 AA.
DE Novel secreted and transmembrane protein PRO1868.
PN US2003017563-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1229
ID ABO25109 standard; protein; 310 AA.
DE Human secreted/transmembrane protein (PRO) #269.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1230
ID ABU64559 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #63.
PN US2002160374-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1231
ID ABU67405 standard; protein; 310 AA.
DE Human secreted protein PRO1868.
PN US2003023054-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1232
ID ABO14925 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1233
ID ABU60813 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #7.
PN US2002160392-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1234
ID ABU67114 standard; protein; 310 AA.
DE Human secreted/transmembrane, PRO, protein SEQ ID 538.
PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1235
ID ABU81236 standard; protein; 310 AA.
DE Human PRO1917polypeptide.
PN US2003032060-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1236
ID ABU69682 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868+H30.
PN US2003017463-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1237
ID ABO14864 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003027143-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1238
ID ADA46057 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1239
ID ADA76488 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1240
ID ADE29627 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003092002-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1241
ID ADA19138 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1242
ID ABO14925 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;

RESULT 1242
ID ADA61761 standard; protein; 310 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1243
ID ADB19546 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1244
ID ADB28087 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1245
ID ADA86566 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1246
ID ADB16130 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087350-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1247
ID ADA47916 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1248
ID ADA18484 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039971-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1249
ID ABO32816 standard; protein; 310 AA.
DE Human secreted/transmembrane protein PRO1868.
PN US2003045693-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1250
ID ADA67711 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1251

ID ADB30718 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1252
ID ADA86014 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082693-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1253
ID ADA97226 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1254
ID ADA79530 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1255
ID ADA87669 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087345-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1256
ID ADB16871 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1257
ID ABO34876 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003044793-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1258
ID ADA16459 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049621-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1259
ID ADA91963 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082694-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1260
ID ADB15026 standard; protein; 310 AA.

DE Human PRO polypeptide #269.
PN US2003087351-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1261
ID ADB18987 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1262
ID ADA94202 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1263
ID ADB20098 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082691-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1264
ID ADB13410 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1265
ID ABO43417 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US200304945-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1266
ID ADA74664 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1267
ID ADA42604 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054401-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1268
ID ADB24897 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077713-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1269
ID ADA82421 standard; protein; 310 AA.
DE Human PRO polypeptide #269.

PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1270
ID ADA75384 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1271
ID ADA85462 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1272
ID ADA84910 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1273
ID ABO17554 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003064367-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1274
ID ADB30166 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1275
ID ADA80694 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1276
ID ADA75936 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1277
ID ADA47161 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1278
ID ADB25457 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077715-A1.

PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1279
ID ADA93633 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US200307721-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1280
ID ADB26983 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1281
ID ADB31270 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1282
ID ABU62957 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US2003054447-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1283
ID ADA61198 standard; protein; 310 AA.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1284
ID ADB24345 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US200307714-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1285
ID ADA96674 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082690-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1286
ID ADA81246 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1287
ID ADA96122 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082759-A1.
PD 01-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1288
ID ADB26431 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1289
ID ADB21916 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1290
ID ADA77695 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068797-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1291
ID ADB18435 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US200307710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1292
ID ADA87118 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1293
ID ADA16883 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039969-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1294
ID ADA13312 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049622-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1295
ID ADA42180 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082540-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1296
ID ADA88221 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1297
ID ADA46609 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1298
ID ADA17527 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003017498-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1299
ID ADA43030 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054351-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1300
ID ADB28639 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1301
ID ADB29191 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1302
ID ABO01894 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003027256-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1303
ID ADA77143 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1304
ID ADA88773 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1305
ID ADA97778 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1306
ID ADB27535 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003022239-A1.
PD 30-JAN-2003.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1307
ID ADB22468 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087344-A1.
PD 08-MAY-2003.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1308
ID ABO17615 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003064923-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1309
ID ADA67159 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1310
ID ADB23020 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003077711-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1311
ID ADB23793 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1312
ID ADA92515 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1313
ID ADB15578 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1314
ID ADB38830 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1315
ID ADB38278 standard; protein; 310 AA.

DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1316
ID ADB66750 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1317
ID ADB98930 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1318
ID ADB90562 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1319
ID ADB77948 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003077654-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1320
ID ADB39663 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1321
ID ADB75084 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082542-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1322
ID ADB47286 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082687-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1323
ID ADB86893 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082697-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1324
ID ADB77498 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.

PN US2003082696-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1325
ID ADB34655 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077717-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1326
ID ADB35759 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1327
ID ADB34103 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077716-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1328
ID ADB35207 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077718-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1329
ID ADB36311 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077720-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1330
ID ADB46706 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082692-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1331
ID ADC28731 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059772-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1332
ID ADC39931 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059828-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1333
ID ADC40445 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059829-A1.

PD 27-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1334
ID ADC19269 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003036061-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1335
ID ADC34569 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003036094-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1336
ID ADC29624 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049676-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1337
ID ADC29155 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049677-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1338
ID ADC41040 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054400-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1339
ID ADC19697 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054441-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1340
ID ADC34145 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003073077-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1341
ID ADC13215 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003073079-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1342
ID AAE38826 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US200307657-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1343
ID ADC50579 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092106-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1344
ID ADC72126 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092107-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1345
ID ADC60105 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092105-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1346
ID ADC53112 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087365-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1347
ID ADC57466 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1348
ID ADC60657 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087367-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1349
ID ADC51132 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1350
ID ADC65659 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1351
ID ADC54757 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.

Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1352
ID ADC53718 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1353
ID ADC59241 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1354
ID ADC56119 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1355
ID ADC58689 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1356
ID ADC2667 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082541-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1357
ID ADC74383 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1016.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1358
ID ADC74606 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1239.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1359
ID ADC74607 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1240.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1360
ID ADD03363 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092104-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1361
ID ADC90355 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1362
ID ADC69774 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194770-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1363
ID ADC48663 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1364
ID ADD10192 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194776-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1365
ID ADD04767 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087354-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1366
ID ADC80723 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1367
ID ADD11230 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1368
ID ADD10551 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1369
ID ADC48111 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1370
ID ADC48111 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.

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RESULT 1370
ID ADD05222 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003104469-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1371
ID ADC80171 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1372
ID ADD11511 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1373
ID ADD09640 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1374
ID ADD04228 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003104381-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1375
ID ADD03804 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003108983-A1.
PD 12-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1376
ID ADD41353 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1377
ID ADD52492 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1378
ID ADD53232 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194792-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1379
ID ADD92672 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1380
ID ADD37304 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1381
ID ADD51940 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1382
ID ADD02739 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1383
ID ADD38106 standard; protein; 310 AA.
DE Human secreted protein #289.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1384
ID ADD38009 standard; protein; 310 AA.
DE Human secreted protein #192.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1385
ID ADD38105 standard; protein; 310 AA.
DE Human secreted protein #288.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1386
ID ADD02173 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1387
ID ADD54355 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1388
ID ADD92672 standard; protein; 310 AA.
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DE Human PRO polypeptide #269.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1389
ID ADE34135 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1390
ID ADE04182 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199057-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1391
ID ADE32479 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1392
ID ADE22411 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1393
ID ADE79635 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1394
ID ADE42171 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1395
ID ADE17988 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1396
ID ADE92120 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1397
ID ADE33583 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003199064-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1398
ID ADE34135 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1399
ID ADE0187 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207417-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1400
ID ADE93224 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194768-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1401
ID ADE19644 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1402
ID ADE35056 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003077583-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1403
ID ADE19092 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1404
ID ADE43288 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1405
ID ADE96077 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
  Query Match      8.2%; Score 138.5; DB 7; Length 310;
  Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1406
ID ADE22963 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199064-A1.
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PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1407
ID ADG23329 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203429-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1408
ID ADE33031 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1409
ID ADE42723 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1410
ID ADD80739 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207418-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1411
ID ADD89767 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1412
ID ADE41051 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1413
ID ADE04850 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1414
ID ADE92979 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1415
ID ADG21688 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207355-A1.
PD 06-NOV-2003.

PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1416
ID ADG23329 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1417
ID ADF97664 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1418
ID ADG80728 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1419
ID ADG80176 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1420
ID ADH62558 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) RAPIER M A.
PA (TUMA/) TOMAS D.
PA (WOOD/) WOOD W I.
Query Match Similarity 21.8%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1421
ID ADH59539 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039972-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1422
ID ADH55468 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1423
ID ADH56020 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1424
ID ADG21688 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207355-A1.
PD 06-NOV-2003.

RESULT 1424
ID ADI38318 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054352-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1425
ID ADI64239 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1426
ID ADI65188 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1427
ID ADI63687 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1428
ID ADH82101 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1429
ID ADH81549 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1430
ID ADJ58518 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003170864-A1.
PD 11-SEP-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1431
ID ADJ26586 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054349-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1432
ID ADM82718 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1433

ID ADN16117 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1434
ID ADN16746 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1435
ID ADN15565 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1436
ID ADN15013 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1437
ID ADC81275 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1438
ID ADE79501 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003135025-A1.
PD 17-JUL-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1439
ID ADD76723 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1440
ID ADD88087 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1441
ID ADD86491 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1442
ID ADE79925 standard; protein; 310 AA.

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DE Human secreted/transmembrane protein, #65.
PN US2003130489-A1.
PD 10-JUL-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1443
ID ADE75939 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1444
ID ADE73601 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003129592-A1.
PD 10-JUL-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1445
ID ADE41512 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1446
ID ADE23515 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1447
ID ADE24067 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1448
ID ADE24710 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1449
ID ADD87535 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1450
ID ADE89401 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1451
ID ADE74136 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211569-A1.
PD 07-AUG-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1452
ID ADE18540 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194794-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1453
ID ADE88849 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199054-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1454
ID ADE99690 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211576-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1455
ID ADE94869 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1456
ID ADE91280 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1457
ID ADE95421 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1458
ID ADE93531 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1459
ID ADF35112 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1460
ID ADE98809 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211569-A1.
```

PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1461
ID ADE92427 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1462
ID ADE90728 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1463
ID ADE91875 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1464
ID ADE99236 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211568-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1465
ID ADG40706 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003225253-A1.
PD 04-DEC-2003.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (MATH/) MATHER J P.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1466
ID ADF74100 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003180312-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1467
ID ADG2454 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1468
ID ADG22240 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1469
ID ADG20310 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1470
ID ADF98216 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1471
ID ADG24433 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1472
ID ADF98787 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1473
ID ADG03618 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1474
ID ADF99339 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1475
ID ADG16924 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1476
ID ADG05383 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1477
ID ADG19650 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;

RESULT 1478
ID ADF73676 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003166051-A1.
PD 04-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1479
ID ADG13487 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1480
ID ADG08544 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1481
ID ADG15714 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1482
ID ADF97112 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1483
ID ADG06297 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1484
ID ADG23881 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1485
ID ADG04170 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1486
ID ADG25071 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207427-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1487
ID ADF73676 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1488
ID ADG07920 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1489
ID ADG55415 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1490
ID ADG61079 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1491
ID ADG62183 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1492
ID ADG92519 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003027145-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1493
ID ADG82384 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1494
ID ADG57623 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207362-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1495
ID ADG57071 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207364-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1496
ID ADG55967 standard; protein; 310 AA.

DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207365-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1497
ID ADG59727 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207368-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1498
ID ADG71093 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207420-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1499
ID ADG92946 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003027146-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1500
ID ADG58175 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207363-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;

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OM protein - protein search, using sw model

Run on: August 26, 2005, 19:28:02 ; Search time 40 Seconds
(without alignments)
772.140 Million cell updates/sec

Title: US-10-767-374-2

Perfect score: 1688

Sequence: 1 MGILLGLLLHLTVDTYGR.....AYIMLCRKTSQEHVYEAAAR 321

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1500 summaries

Database : PIR 79:*

1: pir1: *
2: pir2: *
3: pir3: *
4: pir4: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	178.5	10.6	299	S56749	junctional adhesio
2	172	10.2	7962	I38346	elastic titin - hu
3	169	10.0	365	JC7780	coxsackie- and ade
4	143.5	8.5	1052	B49120	protein-tyrosine k
5	139.5	8.3	3707	S18252	heparan sulfate pr
6	138.5	8.2	1323	PN0568	connectin 3B - chi
7	138.5	8.2	4162	T42633	connectin/titin -
8	136	8.1	584	T08678	hypothetical prote
9	133.5	7.9	5175	T20992	hypothetical prote
10	133.5	7.9	5198	T43290	hemocentin precurs
11	133	7.9	1106	PFHUGB	platelet-derived g
12	132.5	7.8	765	C42632	cell adhesion mole
13	132.5	7.8	812	B42632	cell adhesion mole
14	132.5	7.8	932	A42632	cell adhesion mole
15	132.5	7.8	1272	S26180	neurofascin - chic
16	132	7.8	1896	T088051	Down syndrome cell
17	131	7.8	4391	A38096	perlecan precursor
18	129.5	7.7	1040	A34695	axonal glycoprotei
19	129.5	7.7	3375	T19821	hypothetical prote
20	129	7.6	1427	I51669	tumor suppressor -
21	128.5	7.6	338	JC5519	50K glycoprotein p
22	127.5	7.6	1906	S68235	myosin-light-chain
23	125.5	7.4	1259	A43425	Bravo/Nr-CAM cell
24	125	7.4	351	JH0396	biliary glycoprote
25	124	7.3	853	IJBONC	neural cell adhesi
26	124	7.3	1344	T14316	rig-1 protein - mo
27	123.5	7.3	458	WMMSR1	biliary glycoprote
28	123.5	7.3	521	JC1508	biliary glycoprote
29	123.5	7.3	1091	IJCINHL	neural cell adhesi

30	122.5	7.3	761	1	IJHUNG	neural cell adhesi
31	122.5	7.3	1268	1	A39640	neural cell adhesi
32	120.5	7.1	458	2	JC1509	biliary glycoprote
33	120.5	7.1	538	2	JC2457	vascular cell adhe
34	120	7.1	1033	2	S19247	cell adhesion prot
35	120	7.1	1036	2	S22383	axonin 1 precursor
36	119.5	7.1	417	2	JH0394	biliary glycoprote
37	119.5	7.1	464	2	C30127	transmembrane carc
38	119.5	7.1	483	2	T17346	hypothetical prote
39	119.5	7.1	526	1	A32164	biliary glycoprote
40	119.5	7.1	725	1	IJMSNG	neural cell adhesi
41	119.5	7.1	725	1	JEO100	neural cell adhesi
42	119.5	7.1	1115	1	IJMSNL	neural cell adhesi
43	119	7.0	467	1	HLMSP3	poliovirus recepto
44	119	7.0	1091	2	A58532	glial cell membran
45	119	7.0	28926	1	I38344	titin, cardiac mus
46	118.5	7.0	333	2	A31923	amalgam protein pr
47	118.5	7.0	458	2	S68177	C-CAM2a protein is
48	118.5	7.0	458	2	S23969	cell-adhesion mole
49	118.5	7.0	519	2	A44783	ecto-ATPase precu
50	118.5	7.0	816	2	A49151	fibroblast growth
51	118.5	7.0	1011	2	T13669	neuromusculin - fr
52	118.5	7.0	1694	2	S50065	sialoadhesin - mou
53	118	7.0	547	1	S28904	intercellular adhe
54	118	7.0	1338	2	S09982	protein-tyrosine k
55	117.5	7.0	338	2	JC4776	limbic-system-asso
56	117.5	7.0	344	2	A27681	non-specific cross-
57	117.5	7.0	1092	1	JN0635	neural cell adhesi
58	117.5	7.0	1239	1	A32579	neuroglian - fruit
59	117	6.9	739	2	J80675	vascular cell adhe
60	117	6.9	858	1	IORTNC	neural cell adhesi
61	117	6.9	1447	2	A54100	tumor suppressor p
62	116.5	6.9	806	2	A35963	protein-tyrosine k
63	116.5	6.9	1040	2	A49356	transient axonal g
64	116.5	6.9	1051	2	A39712	kinase-like protei
65	116	6.9	344	2	I56551	neurotrophin rat
66	116	6.9	6658	2	T13931	projectin - fruit
67	115.5	6.8	292	2	S03421	T-cell receptor de
68	115.5	6.8	293	2	A40131	T-cell receptor de
69	115.5	6.8	309	2	I49522	gene B7-2 protein
70	115.5	6.8	725	2	J80099	neural cell adhesi
71	115.5	6.8	738	2	A40096	platelet-endotheli
72	115.5	6.8	1257	1	A41060	neural cell adhesi
73	115	6.8	1209	2	T42718	probable neural ce
74	114.5	6.8	478	2	I53960	PRR2 alpha - human
75	114.5	6.8	521	2	S34338	biliary glycoprote
76	114.5	6.8	538	2	I68093	PRR2 delta - human
77	114.5	6.8	1088	1	IUXLNL	neural cell adhesi
78	114.5	6.8	1173	2	T25893	hypothetical prote
79	114	6.8	544	2	JC5018	intercellular adhe
80	113.5	6.7	530	2	A53437	poliovirus recepto
81	113	6.7	1277	2	T30532	neural cell adhesi
82	112.5	6.7	321	2	JH0395	biliary glycoprote
83	112.5	6.7	345	2	JC4025	opioid-binding cel
84	112.5	6.7	1259	2	S36126	neural cell adhesi
85	112.5	6.7	2029	1	TDFFLK	protein-tyrosine-p
86	112	6.6	946	1	A47299	ror-related recept
87	112	6.6	976	2	T29583	hypothetical prote
88	112	6.6	1273	2	T42405	sax-3 protein - Ca
89	111	6.6	806	1	TVHUF3	fibroblast growth
90	111	6.6	6642	2	T29757	protein UNC-89 - C
91	110.5	6.5	349	2	A34815	carcinoembryonic a
92	110.5	6.5	518	2	JC4024	poliovirus recepto
93	110	6.5	499	2	S33766	platelet-derived g
94	110	6.5	917	2	I48950	telencephalin prec
95	110	6.5	1070	2	JC4593	protein-tyrosine k
96	110	6.5	1880	2	T18531	tractin - medicina
97	110	6.5	2295	2	C88369	protein unc-52 (im
98	109.5	6.5	338	2	JC1238	opioid-binding pro
99	109.5	6.5	345	2	JC1239	opioid-binding pro
100	109.5	6.5	526	2	S70587	butyrophilin precu
101	109.5	6.5	1336	2	I60598	Fit-1 tyrosine kin
102	109	6.5	547	2	B45808	B-lymphocyte antig

833	75	4.4	360	2	AE1931	hypothetical prote	906	74.5	4.4	1446	2	T13018	hypothetical prote
834	75	4.4	362	2	I37120	MHC class I histoc	907	74.5	4.4	1615	2	C75551	glutamate synthase
835	75	4.4	362	2	B30345	MHC class I histoc	908	74.5	4.4	2143	2	G96595	hypothetical prote
836	75	4.4	362	2	I37522	MHC class I histoc	909	74.5	4.4	3890	2	C89921	hypothetical prote
837	75	4.4	362	2	I61904	MHC class I histoc	910	74.5	4.4	4436	2	E71086	hypothetical prote
838	75	4.4	362	2	I54457	MHC class I lympho	911	74	4.4	115	2	A30583	T-cell receptor de
839	75	4.4	362	2	A30345	MHC class I histoc	912	74	4.4	130	2	S08079	Ig kappa chain pre
840	75	4.4	387	2	H65132	hypothetical 44.3	913	74	4.4	143	2	S36321	T-cell receptor de
841	75	4.4	391	1	MHUBT	Ig mu heavy chain	914	74	4.4	145	2	S36299	T-cell receptor de
842	75	4.4	396	2	C95088	S-adenosylmethioni	915	74	4.4	217	2	J80246	Ig lambda chain NI
843	75	4.4	396	2	G97955	methionine adenosy	916	74	4.4	264	2	T26976	hypothetical prote
844	75	4.4	450	1	F0LJFP	gag polyprotein -	917	74	4.4	270	1	S77085	molybdate-binding
845	75	4.4	450	2	S23819	gag protein - feli	918	74	4.4	318	2	C71197	probable ATP-bindi
846	75	4.4	471	2	B38637	Ras inhibitor (clo	919	74	4.4	327	2	F87544	UDP-glucose 4-epim
847	75	4.4	483	2	A69745	hypothetical prote	920	74	4.4	330	2	AE3119	ABC transporter, s
848	75	4.4	496	2	PQ0508	envelope glycoprot	921	74	4.4	341	2	S20827	CAMP response elem
849	75	4.4	496	2	E90181	hypothetical prote	922	74	4.4	351	2	S78042	Ig mu chain C regi
850	75	4.4	507	2	AE0473	colicin (partial)	923	74	4.4	353	2	C86932	conserved hypotnet
851	75	4.4	555	2	T01142	hypothetical prote	924	74	4.4	354	2	S24436	class I histocoma
852	75	4.4	563	2	A70038	L-lactate permease	925	74	4.4	362	2	I56130	HLA-B*5401 - human
853	75	4.4	589	2	A34341	poly(3-hydroxybuty	926	74	4.4	362	2	I59633	MHC HLA-B transmem
854	75	4.4	600	2	A82043	inner membrane cop	927	74	4.4	362	2	I72753	HLA-B*5502 - human
855	75	4.4	630	2	F85074	hypothetical prote	928	74	4.4	362	2	S24434	class I histocoma
856	75	4.4	632	2	S73431	MG288 homolog D09	929	74	4.4	362	2	I72752	HLA-B*5501 - human
857	75	4.4	673	2	S46520	luciferin-binding -	930	74	4.4	362	2	I56133	MHC class I protei
858	75	4.4	686	2	JC7569	Delta-4 protein -	931	74	4.4	365	2	B95260	recf protein limpo
859	75	4.4	808	2	T23129	hypothetical prote	932	74	4.4	365	2	E38125	recombination prot
860	75	4.4	997	2	A40812	Ca2+-transporting	933	74	4.4	387	2	B98168	periplasmic-iron-b
861	75	4.4	997	2	A34307	Ca2+-transporting	934	74	4.4	425	2	E71982	isocitrate dehydro
862	75	4.4	1042	2	B40812	Ca2+-transporting	935	74	4.4	430	2	S50981	probable membrane
863	75	4.4	1234	2	S72640	endo-1,4-beta-xyla	936	74	4.4	430	2	S50981	probable membrane
864	75	4.4	1241	2	S26373	genome polyprotein	937	74	4.4	430	2	S50981	SHS aid-1 (im
865	75	4.4	1277	2	T14152	synaptic scaffold	938	74	4.4	430	2	S50981	SHS aid-1 (im
866	75	4.4	1622	2	AE1717	probable cell surf	939	74	4.4	430	2	S50981	SHS aid-1 (im
867	75	4.4	1806	2	AF1717	probable peptidogl	940	74	4.4	430	2	S50981	SHS aid-1 (im
868	75	4.4	2109	1	I50421	aggreccan precursor	941	74	4.4	430	2	S50981	SHS aid-1 (im
869	75	4.4	2124	2	A28452	proteoglycan core	942	74	4.4	430	2	S50981	SHS aid-1 (im
870	75	4.4	2132	1	A55182	aggreccan precursor	943	74	4.4	430	2	S50981	SHS aid-1 (im
871	75	4.4	2415	1	A33086	aggreccan precursor	944	74	4.4	430	2	S50981	SHS aid-1 (im
872	75	4.4	3216	2	C90538	hypothetical prote	945	74	4.4	430	2	S50981	SHS aid-1 (im
873	75	4.4	3263	2	E82410	hypothetical prote	946	74	4.4	430	2	S50981	SHS aid-1 (im
874	75	4.4	3624	2	AD0835	large repetitive p	947	74	4.4	430	2	S50981	SHS aid-1 (im
875	74.5	4.4	115	2	A24891	T-cell receptor al	948	74	4.4	430	2	S50981	SHS aid-1 (im
876	74.5	4.4	131	2	PQ0059	T-cell receptor be	949	74	4.4	430	2	S50981	SHS aid-1 (im
877	74.5	4.4	133	2	S57885	T-cell receptor al	950	74	4.4	430	2	S50981	SHS aid-1 (im
878	74.5	4.4	135	1	RWHUAV	T-cell receptor al	951	74	4.4	430	2	S50981	SHS aid-1 (im
879	74.5	4.4	142	1	E64794	ybdQ protein - Esc	952	74	4.4	430	2	S50981	SHS aid-1 (im
880	74.5	4.4	142	2	F90709	hypothetical prote	953	74	4.4	430	2	S50981	SHS aid-1 (im
881	74.5	4.4	142	2	B85560	hypothetical prote	954	74	4.4	430	2	S50981	SHS aid-1 (im
882	74.5	4.4	208	2	B49444	Ig lambda chain (N	955	74	4.4	430	2	S50981	SHS aid-1 (im
883	74.5	4.4	212	2	C33258	pregnancy-specific	956	74	4.4	430	2	S50981	SHS aid-1 (im
884	74.5	4.4	235	2	AD1735	hypothetical prote	957	74	4.4	430	2	S50981	SHS aid-1 (im
885	74.5	4.4	294	2	T05381	hypothetical prote	958	74	4.4	430	2	S50981	SHS aid-1 (im
886	74.5	4.4	345	2	C72760	probable neme expo	959	73.5	4.4	430	2	S50981	SHS aid-1 (im
887	74.5	4.4	348	2	I50107	MHC class I histoc	960	73.5	4.4	430	2	S50981	SHS aid-1 (im
888	74.5	4.4	361	2	B90461	hypothetical prote	961	73.5	4.4	430	2	S50981	SHS aid-1 (im
889	74.5	4.4	366	1	HLHWS	MHC class I histoc	962	73.5	4.4	430	2	S50981	SHS aid-1 (im
890	74.5	4.4	377	2	A49885	MHC class I histoc	963	73.5	4.4	430	2	S50981	SHS aid-1 (im
891	74.5	4.4	445	2	S43492	surface antigen -	964	73.5	4.4	430	2	S50981	SHS aid-1 (im
892	74.5	4.4	476	2	A10189	exodeoxyribonuclea	965	73.5	4.4	430	2	S50981	SHS aid-1 (im
893	74.5	4.4	503	2	AB1933	hypothetical prote	966	73.5	4.4	430	2	S50981	SHS aid-1 (im
894	74.5	4.4	625	2	T16777	hypothetical prote	967	73.5	4.4	430	2	S50981	SHS aid-1 (im
895	74.5	4.4	652	2	G96560	hypothetical prote	968	73.5	4.4	430	2	S50981	SHS aid-1 (im
896	74.5	4.4	717	2	B32838	DNA-directed RNA p	969	73.5	4.4	430	2	S50981	SHS aid-1 (im
897	74.5	4.4	732	2	A43315	ETS domain protein	970	73.5	4.4	430	2	S50981	SHS aid-1 (im
898	74.5	4.4	810	1	A33380	interleukin-4 rece	971	73.5	4.4	430	2	S50981	SHS aid-1 (im
899	74.5	4.4	943	2	B45082	neurotrophic recep	972	73.5	4.4	430	2	S50981	SHS aid-1 (im
900	74.5	4.4	946	2	B87316	TonB-dependent rec	973	73.5	4.4	430	2	S50981	SHS aid-1 (im
901	74.5	4.4	1000	2	JE0110	mitotic control pr	974	73.5	4.4	430	2	S50981	SHS aid-1 (im
902	74.5	4.4	1086	2	AH1290	cellobiose-phospho	975	73.5	4.4	430	2	S50981	SHS aid-1 (im
903	74.5	4.4	1302	2	AC2574	hypothetical prote	976	73.5	4.4	430	2	S50981	SHS aid-1 (im
904	74.5	4.4	1333	2	S63403	probable membrane	977	73.5	4.4	430	2	S50981	SHS aid-1 (im
905	74.5	4.4	1356	2	T16718	hypothetical prote	978	73.5	4.4	430	2	S50981	SHS aid-1 (im

979	73.5	4.4	348	2	S09273	Ig alpha chain C r	1052	73	4.3	362	2	184490	lymphocyte antigen
980	73.5	4.4	351	2	B34595	pregnancy-specific	1053	73	4.3	362	2	A45880	MHC class I histoc
981	73.5	4.4	351	2	A97064	D-mannanase hydrol	1054	73	4.3	362	2	154442	MHC class I histoc
982	73.5	4.4	357	2	D82337	UDP-N-acetylenolp	1055	73	4.3	365	2	137482	MHC class I histoc
983	73.5	4.4	362	2	S68090	actin 8 - Arabidop	1056	73	4.3	378	2	S41870	surface antigen -
984	73.5	4.4	366	2	I68712	MHC class I histoc	1057	73	4.3	426	2	C72166	A22R protein - var
985	73.5	4.4	366	2	F72062	hypothetical prote	1058	73	4.3	489	2	T09151	glutathione-disulf
986	73.5	4.4	366	2	A85661	CT449 hypothetical	1059	73	4.3	499	2	E86782	D-alanine activati
987	73.5	4.4	400	2	T34363	hypothetical prote	1060	73	4.3	507	2	A48661	cystathionine beta
988	73.5	4.4	404	2	A46480	FC gamma (IgG) rec	1061	73	4.3	508	2	B91250	hypothetical prote
989	73.5	4.4	408	2	F81252	NADH2 dehydrogenas	1062	73	4.3	521	2	B84746	hypothetical prote
990	73.5	4.4	427	2	F64064	tolB protein - Hae	1063	73	4.3	580	2	A81088	ABC transporter, A
991	73.5	4.4	446	2	S40295	Ig gamma-2a chain	1064	73	4.3	615	2	T20839	hypothetical prote
992	73.5	4.4	451	2	T30603	perlecan homolog 2	1065	73	4.3	681	2	A45055	glutamine-fructose
993	73.5	4.4	482	2	T22754	hypothetical prote	1066	73	4.3	685	2	JC7570	Delta-4 protein -
994	73.5	4.4	551	2	G84301	hypothetical prote	1067	73	4.3	694	2	F97279	TPR-repeat-contain
995	73.5	4.4	554	2	T70512	hypothetical prote	1068	73	4.3	866	2	B85075	probable athila tr
996	73.5	4.4	555	2	D71444	hypothetical prote	1069	73	4.3	885	2	S42841	Ti6G12.1 protein -
997	73.5	4.4	568	2	A89558	acetyl-CoA synthet	1070	73	4.3	913	1	IJCHCR	R-cadherin precurs
998	73.5	4.4	650	2	S22835	alpha-agglutinin -	1071	73	4.3	954	2	A10438	probable exported
999	73.5	4.4	659	2	A85854	hypothetical prote	1072	73	4.3	974	1	URHUAP	peptidylglycine mo
1000	73.5	4.4	659	2	G91009	colicin I receptor	1073	73	4.3	977	2	F45877	protein-tyrosine k
1001	73.5	4.4	663	1	QREIC	colicin I receptor	1074	73	4.3	998	2	H75005	ATP-dependent prot
1002	73.5	4.4	666	2	D82511	1,4-alpha-glucan b	1075	73	4.3	1071	2	B84062	hypothetical prote
1003	73.5	4.4	721	2	T05815	hypothetical prote	1076	73	4.3	1115	2	T29012	hypothetical prote
1004	73.5	4.4	737	2	T46243	hypothetical prote	1077	73	4.3	1137	2	E86708	pyruvate carboxyla
1005	73.5	4.4	743	2	D64062	GTP diphosphokinas	1078	73	4.3	1144	2	A81983	probable DNA-direc
1006	73.5	4.4	780	2	A34102	von Willebrand fac	1079	73	4.3	1177	2	T16594	hypothetical prote
1007	73.5	4.4	785	2	S54016	SOR2 protein - yea	1080	73	4.3	1199	2	T23005	hypothetical prote
1008	73.5	4.4	790	2	I51638	F-cadherin - AfriC	1081	73	4.3	1289	2	C70044	probable phosphoe
1009	73.5	4.4	794	2	T36972	probable membrane	1082	73	4.3	1293	2	E85557	enterobactin synth
1010	73.5	4.4	813	1	A49123	fibroblast growth	1083	73	4.3	1293	2	A90707	enterobactin synth
1011	73.5	4.4	868	2	D86349	hypothetical prote	1084	73	4.3	1311	2	T33757	hypothetical prote
1012	73.5	4.4	886	2	E75625	hypothetical prote	1085	73	4.3	1407	1	T00558	probable ABC trans
1013	73.5	4.4	891	2	T19915	hypothetical prote	1086	73	4.3	1437	2	T31093	probable protein-t
1014	73.5	4.4	899	1	GNVMVM	pol polyprotein -	1087	73	4.3	1452	1	S17669	protein-tyrosine-p
1015	73.5	4.4	1117	1	S33851	fibronectin-bindin	1088	73	4.3	1484	2	T42632	breast cancer tumo
1016	73.5	4.4	1127	1	E71156	endopeptidase La h	1089	73	4.3	1615	2	B49502	protein-tyrosine-p
1017	73.5	4.4	1133	1	GNVUSR	M polyprotein prec	1090	73	4.3	1714	1	S18644	multifunctional am
1018	73.5	4.4	1133	2	S12597	M polyprotein prec	1091	73	4.3	1767	2	A49502	protein-tyrosine-p
1019	73.5	4.4	1166	2	T28680	fibrogen-binding	1092	73	4.3	2118	2	S72705	mycoserolate synth
1020	73.5	4.4	1259	2	H65233	ytfn protein - Esc	1093	73	4.3	2505	1	XYRTEA	enoyl-lacyl-carrie
1021	73.5	4.4	1293	1	YGECEF	enterobactin synth	1094	73	4.3	2550	2	B53435	vesicular transport
1022	73.5	4.4	1507	2	D97106	large chain of NAD	1095	73	4.3	2824	2	T22759	hypothetical prote
1023	73.5	4.4	2218	2	B84683	hypothetical prote	1096	73	4.3	3947	2	T52486	ferrichrome sidero
1024	73.5	4.4	2761	2	T21064	hypothetical prote	1097	73	4.3	15281	2	S41309	cyclosporin synthe
1025	73.5	4.4	2899	2	T21546	hypothetical prote	1098	73	4.3	104	2	S36064	Ig lambda chain -
1026	73.5	4.4	2915	2	G87867	protein F36A2.13 [1099	72.5	4.3	110	2	S23368	T-cell receptor al
1027	73.5	4.4	3097	2	T28635	glutamate synthase	1100	72.5	4.3	110	2	S22897	T-cell receptor al
1028	73	4.3	107	2	B45722	anti-glycoprotein	1101	72.5	4.3	111	1	L6HULT	Ig lambda chain V-
1029	73	4.3	113	2	I46637	rearranged T-cell	1102	72.5	4.3	112	1	L1HURA	Ig lambda chain V-
1030	73	4.3	113	2	B49041	T-cell receptor al	1103	72.5	4.3	114	2	T38315	T-cell receptor be
1031	73	4.3	120	2	B25429	T-cell receptor be	1104	72.5	4.3	114	2	T38314	T-cell receptor be
1032	73	4.3	126	2	B46538	Ig heavy chain, me	1105	72.5	4.3	115	2	S03511	T-cell receptor be
1033	73	4.3	131	1	L6HUEB	Ig lambda chain pr	1106	72.5	4.3	115	2	S03510	T-cell receptor be
1034	73	4.3	135	2	S00388	T-cell receptor ga	1107	72.5	4.3	133	2	A25777	T-cell receptor be
1035	73	4.3	162	2	E71131	hypothetical prote	1108	72.5	4.3	135	1	RWHUVY	T-cell receptor CK
1036	73	4.3	173	2	T27373	peptidylprolyl too	1109	72.5	4.3	135	2	S57877	rearranged t-cell
1037	73	4.3	175	2	I38408	neu differentiation	1110	72.5	4.3	136	2	I46635	Ig lambda chain pr
1038	73	4.3	186	2	I61783	sodium channel bet	1111	72.5	4.3	140	2	PH0132	hypothetical prote
1039	73	4.3	210	2	I49294	CD7 antigen - mous	1112	72.5	4.3	224	2	B81783	42K surface glycop
1040	73	4.3	213	2	S21066	Ig lambda chain V	1113	72.5	4.3	233	2	JH0372	Ig heavy chain VH1
1041	73	4.3	244	2	S12328	Ig heavy chain C r	1114	72.5	4.3	249	2	S69340	hypothetical prote
1042	73	4.3	247	2	AF0869	probable fimbrial	1115	72.5	4.3	250	2	D83835	T-cell receptor be
1043	73	4.3	251	2	T15495	hypothetical prote	1116	72.5	4.3	307	1	RWMSBC	hypothetical prote
1044	73	4.3	253	2	G97267	PHP superfamily hy	1117	72.5	4.3	312	2	T33344	UDP-glucose 4-epim
1045	73	4.3	267	1	RWMSCH	T-cell receptor al	1118	72.5	4.3	334	2	D83788	class I histocompa
1046	73	4.3	354	2	E82850	fimbrial adhesin p	1119	72.5	4.3	365	2	JH0537	MHC class I histoc
1047	73	4.3	362	1	HLHUB8	MHC class I histoc	1120	72.5	4.3	366	2	I54430	MHC class I histoc
1048	73	4.3	362	2	A45834	MHC class I histoc	1121	72.5	4.3	366	2	I61866	MHC HLA-Cw2.2 chai
1049	73	4.3	362	2	I61907	MHC class I histoc	1122	72.5	4.3	366	2	I56034	gene HLA-C protein
1050	73	4.3	362	2	A45850	MHC class I histoc	1123	72.5	4.3	408	2	S76830	hypothetical prote
1051	73	4.3	362	2	I81233	lymphocyte antigen	1124	72.5	4.3	416	1	KIVKGL	phosphoglycerate k

1125	72.5	4.3	426	2	D42519	A20R protein - vac	1198	72	4.3	997	2	S23444	Ca2+-transporting
1126	72.5	4.3	426	2	T37408	probable 49.1k pro	1199	72	4.3	997	2	S04651	Ca2+-transporting
1127	72.5	4.3	432	2	T43476	hypothetical prote	1200	72	4.3	997	2	B31982	Ca2+-transporting
1128	72.5	4.3	460	1	T38608	hypothetical prote	1201	72	4.3	1042	1	PWRBMC	Ca2+-transporting
1129	72.5	4.3	468	1	VGBEEH	glycoprotein gp13	1202	72	4.3	1042	2	A31981	Ca2+-transporting
1130	72.5	4.3	471	1	PASCA	alkaline phosphata	1203	72	4.3	1042	2	S04652	Ca2+-transporting
1131	72.5	4.3	481	2	S62427	G-protein signalin	1204	72	4.3	1042	2	A33881	Ca2+-transporting
1132	72.5	4.3	491	1	IJBOPC	P-cadherin - bovin	1205	72	4.3	1043	2	A31982	Ca2+-transporting
1133	72.5	4.3	502	2	A83938	lipopolysaccharide	1206	72	4.3	1135	1	GNVUHV	M polyprotein - Ha
1134	72.5	4.3	503	2	JCS287	SHP substrate-1 pr	1207	72	4.3	1135	1	GNVUHV	M polyprotein - Ha
1135	72.5	4.3	533	1	T34458	hypothetical prote	1208	72	4.3	1218	2	A00837	probable ABC trans
1136	72.5	4.3	553	1	H46329	cell fusion glycop	1209	72	4.3	1218	2	T30293	ABC transport prot
1137	72.5	4.3	568	2	S57830	glucose-6-phosphat	1210	72	4.3	1289	2	E90098	RNA polymerase III
1138	72.5	4.3	569	2	S41806	glucose-6-phosphat	1211	72	4.3	1306	2	S25370	MSB2 protein - yea
1139	72.5	4.3	612	2	B48936	RNA polymerase sig	1212	72	4.3	1408	2	H69068	cell surface glyco
1140	72.5	4.3	619	2	A43361	Ets-related transa	1213	72	4.3	1695	2	A56921	kinesin family pro
1141	72.5	4.3	641	2	G85043	hypothetical prote	1214	72	4.3	2090	2	S26058	probable transform
1142	72.5	4.3	694	2	G01161	thymopoietin alpa	1215	72	4.3	2142	1	ZLVNPF	genome polyprotein
1143	72.5	4.3	723	2	T30094	hypothetical prote	1216	72	4.3	2302	2	T14328	protein-tyrosine-p
1144	72.5	4.3	780	1	S43859	ATPase - Sulfolobu	1217	72	4.3	2492	1	NNWVTD	nonstructural poly
1145	72.5	4.3	820	2	T14879	hypothetical prote	1218	72	4.3	5188	2	B85547	probable RTX famil
1146	72.5	4.3	867	2	AD1856	hypothetical prote	1219	72	4.3	5291	2	F90696	hypothetical prote
1147	72.5	4.3	899	2	B75018	hypothetical prote	1220	71.5	4.2	101	2	D25733	T-cell receptor al
1148	72.5	4.3	923	2	S09583	peptidylglycine mo	1221	71.5	4.2	102	2	S29588	Ig kappa chain V r
1149	72.5	4.3	976	1	TVHUKT	protein-tyrosine k	1222	71.5	4.2	112	2	S22891	T-cell receptor al
1150	72.5	4.3	980	2	A38523	genome polyprotein	1223	71.5	4.2	113	2	S03410	Ig kappa chain pre
1151	72.5	4.3	1088	1	PFRTGA	platelet-derived g	1224	71.5	4.2	139	1	RWHU7A	T-cell receptor al
1152	72.5	4.3	1131	2	T15787	hypothetical prote	1225	71.5	4.2	142	2	S36310	T-cell receptor de
1153	72.5	4.3	1143	2	S46122	SNF2 protein homol	1226	71.5	4.2	148	2	A32536	T-cell receptor al
1154	72.5	4.3	1663	1	C3MS	complement C3 prec	1227	71.5	4.2	155	2	S81878	T cell antigen rec
1155	72.5	4.3	1707	2	AH0095	two-component hybr	1228	71.5	4.2	155	2	S71257	major latex protei
1156	72.5	4.3	2201	2	AD0095	probable sideropho	1229	71.5	4.2	246	2	PC4397	mucin 3 T10 - huma
1157	72.5	4.3	3283	2	AC1018	large repetitive p	1230	71.5	4.2	255	1	S48146	mucin 1 precursor,
1158	72	4.3	111	2	S26256	T-cell receptor be	1231	71.5	4.2	259	2	S60617	hypothetical prote
1159	72	4.3	111	2	S28255	T-cell receptor be	1232	71.5	4.2	270	2	G84163	hypothetical prote
1160	72	4.3	111	2	S09963	Ig kappa chain V-J	1233	71.5	4.2	291	2	JQ1562	hypothetical 33.9K
1161	72	4.3	136	2	S36304	T-cell receptor de	1234	71.5	4.2	299	2	I46690	CD80 precursor - r
1162	72	4.3	133	2	PN0538	Ig heavy chain V r	1235	71.5	4.2	329	2	JCS168	UDFglucose 4-epime
1163	72	4.3	144	2	A27577	T-cell receptor al	1236	71.5	4.2	337	2	S65022	glucan endo-1,3-be
1164	72	4.3	171	2	S38237	hypothetical prote	1237	71.5	4.2	345	2	I68749	MHC class I lympo
1165	72	4.3	259	2	F69678	involved in polyke	1238	71.5	4.2	352	2	G81921	hypothetical prote
1166	72	4.3	281	1	S34626	translation elonga	1239	71.5	4.2	364	2	G44086	glycerophosphodie
1167	72	4.3	285	2	S36903	Fc gamma (IgG) rec	1240	71.5	4.2	365	1	S76914	translation releas
1168	72	4.3	296	2	I46021	PC gamma receptor	1241	71.5	4.2	366	2	I37526	MHC class I histoc
1169	72	4.3	323	1	PEPLBJ	penicillopepsin (B	1242	71.5	4.2	366	2	I72113	MHC histocompatibi
1170	72	4.3	333	2	JC7713	ankyrin-repeat pro	1243	71.5	4.2	366	2	I59622	lymphocyte antigen
1171	72	4.3	336	2	I49582	CD1.1 - mouse	1244	71.5	4.2	366	2	I38507	MHC class I histoc
1172	72	4.3	350	2	I54308	MHC HLA B*1 - huma	1245	71.5	4.2	370	2	A39115	glucan endo-1,3-be
1173	72	4.3	354	2	S24437	class I histocompa	1246	71.5	4.2	380	2	E87729	protein Y23HSA.4 [
1174	72	4.3	362	2	S24435	class I histocompa	1247	71.5	4.2	398	1	S45545	GTP cyclohydrolase
1175	72	4.3	362	2	S16789	class I histocompa	1248	71.5	4.2	398	2	H96967	protein containing
1176	72	4.3	365	2	I72170	MHC class I histoc	1249	71.5	4.2	426	2	T28563	hypothetical prote
1177	72	4.3	365	2	I38610	MHC class I histoc	1250	71.5	4.2	426	2	C36850	A21R protein - var
1178	72	4.3	369	2	S12406	glucan endo-1,3-be	1251	71.5	4.2	428	1	AUR2QD	glutamate-ammonia
1179	72	4.3	386	2	A41950	retrovirus-related	1252	71.5	4.2	436	2	A53568	methylocytoxyl-Co
1180	72	4.3	397	2	B87343	conserved hypotet	1253	71.5	4.2	468	1	B84540	acid phosphatase (
1181	72	4.3	406	2	B35878	class I major hist	1254	71.5	4.2	492	2	T47720	pyruvate kinase-li
1182	72	4.3	416	1	A42879	advanced glycosyla	1255	71.5	4.2	497	2	E86485	hypothetical prote
1183	72	4.3	436	2	T16638	hypothetical prote	1256	71.5	4.2	528	2	PC4025	intercellular adhe
1184	72	4.3	450	1	MHDG	Ig mu chain C regi	1257	71.5	4.2	539	2	S16989	dihydroliipoamide S
1185	72	4.3	456	2	T38221	hypothetical serin	1258	71.5	4.2	567	2	AF0274	ribulokinase (EC 2
1186	72	4.3	505	2	T07883	cellulase (EC 3.2.	1259	71.5	4.2	586	2	T19075	hypothetical prote
1187	72	4.3	520	2	A71564	hypothetical prote	1260	71.5	4.2	591	2	A80509	oxaloacetate decar
1188	72	4.3	569	2	A36187	interleukin-1 rece	1261	71.5	4.2	591	2	A50909	oxaloacetate decar
1189	72	4.3	580	2	T28725	hypothetical prote	1262	71.5	4.2	640	1	A55073	transforming prote
1190	72	4.3	592	2	T43402	probable protein k	1263	71.5	4.2	668	1	A42908	meprin A (EC 3.4.2
1191	72	4.3	822	1	IJWSCP	P-cadherin precurs	1264	71.5	4.2	687	2	T09051	PePA protein - Pse
1192	72	4.3	828	2	T22367	hypothetical prote	1265	71.5	4.2	702	2	B69498	hypothetical prote
1193	72	4.3	915	2	T23937	hypothetical prote	1266	71.5	4.2	704	2	I39805	cyclomaltodextrin
1194	72	4.3	974	1	A49714	protein-tyrosine k	1267	71.5	4.2	710	2	A96540	hypothetical prote
1195	72	4.3	994	2	A32792	Ca2+-transporting	1268	71.5	4.2	710	2	B71417	hypothetical prote
1196	72	4.3	997	1	PWRBSC	Ca2+-transporting	1269	71.5	4.2	747	2	B47093	cellulase (EC 3.2.
1197	72	4.3	997	2	B31981	Ca2+-transporting	1270	71.5	4.2	761	1	TVHUMB	transforming prote

1271	71.5	4.2	811	2	PN0689	connectin 1 - chic	1344	71	4.2	1330	2	B70836	hypothetical prote
1272	71.5	4.2	825	2	A59296	alpha-L-arabinofur	1345	71	4.2	1554	2	C72647	hypothetical prote
1273	71.5	4.2	828	2	S52393	beta-galactosidase	1346	71	4.2	1653	2	B91052	hypothetical prote
1274	71.5	4.2	857	2	T04208	probable anthranil	1347	71	4.2	1653	2	F85896	hypothetical prote
1275	71.5	4.2	960	1	BN0677	protein-tyrosine k	1348	71	4.2	2256	2	AD1018	large repetitive p
1276	71.5	4.2	1137	2	B90734	probable host spec	1349	71	4.2	4351	2	T00252	MEGF1 protein - r
1277	71.5	4.2	1138	2	D85584	probable tail comp	1350	70.5	4.2	98	2	PH1061	ig light chain v r
1278	71.5	4.2	1179	2	C36792	hypothetical prote	1351	70.5	4.2	107	2	A26945	T-cell receptor de
1279	71.5	4.2	1211	2	S68251	phospholipase C, i	1352	70.5	4.2	108	2	C30502	ig kappa chain v r
1280	71.5	4.2	1226	2	JC7503	protein-tyrosine p	1353	70.5	4.2	115	2	B26524	T-cell receptor be
1281	71.5	4.2	1345	2	S55669	tegument protein 7	1354	70.5	4.2	115	2	A30995	T-cell receptor be
1282	71.5	4.2	1441	2	A86685	prophage p11 prote	1355	70.5	4.2	128	2	PN0445	ig kappa chain pre
1283	71.5	4.2	1490	2	JC5145	DNA (cytosine-5)-	1356	70.5	4.2	137	2	S03477	T-cell receptor al
1284	71.5	4.2	1512	2	AB1347	probable peptidogl	1357	70.5	4.2	142	2	TI0577	conserved hypothet
1285	71.5	4.2	1630	2	A53577	ascites stialoglyco	1358	70.5	4.2	180	2	TI8313	hypothetical prote
1286	71.5	4.2	1815	2	B95942	conserved hypothet	1359	70.5	4.2	225	2	AE2347	ribonuclease Hrl l
1287	71.5	4.2	1939	2	D97316	probable S-layer p	1360	70.5	4.2	238	2	S29575	ig light chain - r
1288	71.5	4.2	1946	2	AC2141	serine/threonine k	1361	70.5	4.2	235	2	S25754	ig lambda chain - r
1289	71.5	4.2	3131	2	S39842	enniatin synthetas	1362	70.5	4.2	266	2	B82655	UPP-3-O-(R-3-hydro
1290	71.5	4.2	3191	2	T22945	hypothetical prote	1363	70.5	4.2	269	2	H84313	hypothetical prote
1291	71	4.2	87	2	PH1082	ig light chain v r	1364	70.5	4.2	272	2	C75560	conserved hypothet
1292	71	4.2	98	2	S28911	ig heavy chain v r	1365	70.5	4.2	292	2	C84461	En/Spm-like transp
1293	71	4.2	115	1	KWMSL7	ig kappa chain pre	1366	70.5	4.2	328	2	S65023	glucan endo-1,3-be
1294	71	4.2	166	2	PL0012	ig heavy chain pre	1367	70.5	4.2	340	2	A95939	probable spermidin
1295	71	4.2	180	2	G69222	molybdenum formylm	1368	70.5	4.2	345	2	F82528	hypothetical prote
1296	71	4.2	254	2	S48547	probable membrane	1369	70.5	4.2	348	2	S09270	ig alpha chain C r
1297	71	4.2	287	2	E69901	cell wall-binding	1370	70.5	4.2	355	2	B97795	peptide chain rele
1298	71	4.2	321	2	AC0821	probable exported	1371	70.5	4.2	372	2	T09962	cyclin A-type - Ma
1299	71	4.2	336	2	C27658	pregnancy-specific	1372	70.5	4.2	389	2	A44832	esterase estA - Ps
1300	71	4.2	357	2	S09269	ig alpha chain C r	1373	70.5	4.2	397	2	S52783	aspartic proteinas
1301	71	4.2	362	2	JH0541	class I histocompa	1374	70.5	4.2	406	2	D90492	hypothetical prote
1302	71	4.2	362	2	JH0539	class I histocompa	1375	70.5	4.2	419	2	A36109	pregnancy-specific
1303	71	4.2	362	2	JH0540	class I histocompa	1376	70.5	4.2	424	1	A36000	sperm-binding glyc
1304	71	4.2	362	2	I56149	lymphocyte antigen	1377	70.5	4.2	448	2	A84410	hydroxymethylpyrim
1305	71	4.2	362	2	I84488	lymphocyte antigen	1378	70.5	4.2	445	2	S56260	probable membrane
1306	71	4.2	362	2	I54314	MHC HLA-B39N - hum	1379	70.5	4.2	456	2	C57742	cyclin II - maize
1307	71	4.2	362	2	I54505	MHC HLA-B39N - hum	1380	70.5	4.2	482	2	JB0395	phospho-beta-galac
1308	71	4.2	362	2	I59645	HLA-B-6701 - human	1381	70.5	4.2	483	2	E71681	isocitrate dehydro
1309	71	4.2	362	2	I54298	gene HLA-B protein	1382	70.5	4.2	485	2	S36772	E-selectin - bovin
1310	71	4.2	362	2	I68850	MHC class I histoc	1383	70.5	4.2	493	2	S07375	flagellin H-1 - Sa
1311	71	4.2	364	2	S35997	MHC class I histoc	1384	70.5	4.2	520	2	A64033	hypothetical prote
1312	71	4.2	367	1	MHCH	ig mu chain C regi	1385	70.5	4.2	532	1	A29849	intercellular adhe
1313	71	4.2	368	2	D90607	hypothetical prote	1386	70.5	4.2	535	2	JH0263	carboxy-terminal p
1314	71	4.2	374	2	F69323	carbamoyl-phosphat	1387	70.5	4.2	559	2	T43847	DNA-directed RNA p
1315	71	4.2	425	2	C64523	isocitrate dehydro	1388	70.5	4.2	577	2	A95902	probable glycerol-
1316	71	4.2	427	2	T05019	hypothetical prote	1389	70.5	4.2	600	2	AH2623	60 kd inner-membra
1317	71	4.2	427	2	A11264	N-acetylmuramoyl-L	1390	70.5	4.2	600	2	G97405	60K inner-membra
1318	71	4.2	453	2	H96798	hypothetical prote	1391	70.5	4.2	612	2	T13152	WDRI protein - hum
1319	71	4.2	459	2	T43538	zinc finger protei	1392	70.5	4.2	629	2	AF0079	RNA polymerase sig
1320	71	4.2	476	2	A70318	aldehyde dehydroge	1393	70.5	4.2	629	2	A46500	Ly-9.2 antigen - m
1321	71	4.2	481	2	QJ1147	N-acetylmuramoyl-L	1394	70.5	4.2	666	2	C90464	hypothetical prote
1322	71	4.2	509	2	AC2217	hypothetical prote	1395	70.5	4.2	682	2	T10319	envelope protein E
1323	71	4.2	511	1	VGBEF4	glycoprotein C - h	1396	70.5	4.2	727	2	D75122	hypothetical prote
1324	71	4.2	513	2	A48233	polymavirus enhan	1397	70.5	4.2	764	2	T39194	5-methyltetrahydro
1325	71	4.2	550	2	G70597	probable proteinas	1398	70.5	4.2	774	2	A39832	scabrous locus (sc
1326	71	4.2	626	2	S53871	PmeI 17 protein -	1399	70.5	4.2	782	2	AF0179	conserved hypothet
1327	71	4.2	656	2	D96831	hypothetical prote	1400	70.5	4.2	787	2	A75347	GTP pyrophosphokin
1328	71	4.2	662	2	JC7906	sucrose 1f-fructos	1401	70.5	4.2	846	1	PNECA	penicillin amidase
1329	71	4.2	681	2	I53743	glutamine-fructose	1402	70.5	4.2	856	2	I58411	protein-tyrosine k
1330	71	4.2	744	2	AF0410	GTP diphosphokinas	1403	70.5	4.2	874	2	E97302	hypothetical prote
1331	71	4.2	771	2	T13618	hypothetical prote	1404	70.5	4.2	878	2	AD0664	probable exported
1332	71	4.2	776	2	B41704	genome polypotein	1405	70.5	4.2	879	2	S64755	hypothetical prote
1333	71	4.2	776	2	A41704	genome polypotein	1406	70.5	4.2	905	1	IJXLQ1	N-cadherin 1 precu
1334	71	4.2	796	2	A90541	hypothetical prote	1407	70.5	4.2	1043	2	F97302	hypothetical prote
1335	71	4.2	972	1	TVHMD	macrophage colony-	1408	70.5	4.2	1133	1	A43964	M polypotein prec
1336	71	4.2	976	1	TVMSMD	macrophage colony-	1409	70.5	4.2	1134	1	A43960	M polypotein prec
1337	71	4.2	994	2	A49849	Ca2+-transporting	1410	70.5	4.2	1134	1	GNVU22	M polypotein prec
1338	71	4.2	994	2	A70776	probable glng - My	1411	70.5	4.2	1210	2	S35548	DNA-directed RNA p
1339	71	4.2	1001	1	PWRBFC	Ca2+-transporting	1412	70.5	4.2	1603	1	VJWK5	vitellogenin vit-5
1340	71	4.2	1067	2	TJ8663	hypothetical prote	1413	70.5	4.2	1617	2	B86483	protein F5J5.15 [i
1341	71	4.2	1135	2	JS0605	M polypotein - Ha	1414	70.5	4.2	1666	1	C3GP	complement C3 prec
1342	71	4.2	1142	1	GNVUPH	M polypotein prec	1415	70.5	4.2	1804	2	AI1850	serine/threonine k
1343	71	4.2	1157	2	F97255	fusion of alpha-gl	1416	70.5	4.2	2037	2	TI6881	hypothetical prote

1417 70.5 4.2 3097 2 T00021 DN-cadherin - frui
1418 70.5 4.2 3898 1 GNVVHB genome polyprotein
1419 70 4.1 110 2 B30583 T-cell receptor de
1420 70 4.1 118 2 B32536 T-cell receptor al
1421 70 4.1 132 2 PL0114 Ig lambda chain pr
1422 70 4.1 132 2 S04937 Ig lambda chain pr
1423 70 4.1 132 2 S05268 Ig kappa chain pre
1424 70 4.1 133 2 S57886 T cell receptor al
1425 70 4.1 133 2 B30587 outer membrane pro
1426 70 4.1 135 2 JQ0472 T-cell receptor be
1427 70 4.1 136 2 B45893 T-cell receptor al
1428 70 4.1 224 2 C72390 hypothetical prote
1429 70 4.1 233 2 S25752 Ig lambda chain -
1430 70 4.1 239 2 T48937 hypothetical prote
1431 70 4.1 248 2 PQ0769 glycoprotein G - b
1432 70 4.1 261 2 S29360 Fc gamma (IgG) rec
1433 70 4.1 263 2 JQ2284 glycoprotein G - b
1434 70 4.1 275 1 HLHU10 MHC class I histoc
1435 70 4.1 276 2 T47734 hypothetical prote
1436 70 4.1 294 2 H90300 hypothetical prote
1437 70 4.1 295 2 C83231 conserved hypothet
1438 70 4.1 303 1 B36227 urate oxidase (EC
1439 70 4.1 323 2 T50255 hypothetical trans
1440 70 4.1 327 2 JT0584 deoxyribonuclease
1441 70 4.1 354 2 I59308 class I histocompa
1442 70 4.1 354 2 S24440 MHC class I histoc
1443 70 4.1 361 2 I54418 MHC class I histoc
1444 70 4.1 362 2 I37520 MHC class I histoc
1445 70 4.1 362 2 I61860 MHC HLA-B*8 chain
1446 70 4.1 368 2 I55961 MHC class I histoc
1447 70 4.1 373 2 T38687 hypothetical prote
1448 70 4.1 374 2 T46065 hypothetical prote
1449 70 4.1 380 1 S53307 alcohol dehydrogen
1450 70 4.1 382 1 A70078 conserved hypothet
1451 70 4.1 385 2 G71246 hypothetical prote
1452 70 4.1 392 2 D83513 probable esterase
1453 70 4.1 400 1 SAVLA large surface anti
1454 70 4.1 446 2 T41091 conserved hypothet
1455 70 4.1 452 2 S77538 serine proteinase
1456 70 4.1 459 2 T37704 zinc-finger protei
1457 70 4.1 485 2 AD0041 rhamnulokinase (EC
1458 70 4.1 492 2 F81823 conserved hypothet
1459 70 4.1 515 2 JC7533 inulinase (EC 3.2.
1460 70 4.1 517 2 T02464 hypothetical prote
1461 70 4.1 557 2 H69678 involved in polyke
1462 70 4.1 564 2 AH1696 alpha-acetolactate
1463 70 4.1 570 2 A57535 intrileukin 1 recep
1464 70 4.1 570 2 A75201 hypothetical prote
1465 70 4.1 580 2 AE1452 ABC transporter, A
1466 70 4.1 619 2 G72709 probable DNA ligas
1467 70 4.1 622 2 A61197 6-methylsalilic
1468 70 4.1 625 2 G70574 dnaK-type molecula
1469 70 4.1 632 2 T00108 hypothetical prote
1470 70 4.1 634 2 T51282 beta-D-glucan exoh
1471 70 4.1 664 2 A84716 probable GTP-bindi
1472 70 4.1 704 2 F87706 prolly oligopeptid
1473 70 4.1 729 1 A60006 coat protein vp1 -
1474 70 4.1 753 2 F90933 catalase HP11 [imp
1475 70 4.1 753 2 B85782 catalase, hydropor
1476 70 4.1 760 2 T24521 hypothetical prote
1477 70 4.1 804 2 S64090 SCV1 protein - yea
1478 70 4.1 814 2 T00740 hypothetical prote
1479 70 4.1 817 2 A48721 titin, muscle - ch
1480 70 4.1 848 2 C70834 probable endopepti
1481 70 4.1 883 2 S57653 brevicain precursor
1482 70 4.1 883 2 S49126 hypothetical prote
1483 70 4.1 918 2 T02759 translation initia
1484 70 4.1 952 2 S64473 male-specific leth
1485 70 4.1 955 2 S52959 hypothetical prote
1486 70 4.1 963 2 AF2119 receptor-protein t
1487 70 4.1 975 2 I48974 protein-tyrosine k
1488 70 4.1 978 1 A49814 regulatory factor
1489 70 4.1 979 2 A35913

1490 70 4.1 988 2 T08102 myrosinase-binding
1491 70 4.1 1001 2 T00532 probable cadmium-t
1492 70 4.1 1016 2 T30943 aminopeptidase (EC
1493 70 4.1 1025 2 A83186 probable RND efflu
1494 70 4.1 1089 1 S33727 platelet-derived g
1495 70 4.1 1108 1 S45917 platelet serine/th
1496 70 4.1 1133 2 E86308 hypothetical prote
1497 70 4.1 1172 2 F84572 probable cadmium-t
1498 70 4.1 1189 2 T17088 homeodomain-intera
1499 70 4.1 1402 2 D70634 probable polyketid
1500 70 4.1 1409 2 T42522 protein-tyrosine-p

ALIGNMENTS

RESULT 1

S56749
Junctional adhesion molecule precursor - human
N/Alternate names: Fli platelet antigen; platelet adhesion molecule PAM-1; platelet Fli
C/Species: Homo sapiens (man)
C/Date: 27-Oct-1995 #sequence_revision 01-Feb-2002 #text_change 09-Jul-2004
C/Accession: A59406; S56749
R/Ozaki, H.; Ishii, K.; Horiuchi, H.; Arai, H.; Kawamoto, T.; Okawa, K.; Iwamatsu, A.; K
J. Immunol. 163, 553-557, 1999
A/Title: Cutting edge: combined treatment of TNF-alpha and IFN-gamma causes redistributi
A/Reference number: A59406; MUID:99323940; PMID:10395639
A/Accession: A59406
A/Status: preliminary
A/Molecule type: DNA
A/Residues: 1-299 <OZA>
A/Cross-references: UNIPROT:Q9V624; GB:AAD42050; NID:G5326797; PIDN:AAD42050.1
R/Naik, U.P.; Ehrlich, Y.H.; Kornecki, E.
Biochem. J. 310, 155-162, 1995
A/Title: Mechanisms of platelet activation by a stimulatory antibody: cross-linking of a
A/Reference number: S56749; MUID:95374438; PMID:7646439
A/Accession: S56749
A/Molecule type: protein
A/Residues: 28-49 'X', 51-53:62-73 'E', 75-103:123 'F', 125-130 'FDKDXITLYNKY', 'LT', 206, 'X',
A/Note: the order of the peptides other than the amino terminus was not determined
C/Genetics:
A/Gene: JAM
C/Keywords: glycoprotein; phosphoprotein; platelet aggregation; platelet membrane
F/1-25/Domain: signal sequence #status predicted <SIG>
F/26-299/Product: junctional adhesion molecule #status predicted <MAT>

Query Match 10.6%; Score 178.5; DB 2; Length 299;

Best Local Similarity 26.5%; Pred. No. 2.3e-06;

Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGILLGLLLGHLTVDTYGRPILEVPSVTGPKGDNVNLCTYDPLOQYTVLVKWLQVR 60

Db 17 LAILLCSALGSGVTVHS--SEFEVRIPEN-----NPVKLSLAY---SGFSRVRW---- 62

QY 61 GSDPVTIFLRDSSGDHIQAKYQGRHLVSHK-----VPGDVSLOQLSTLEMDRSHYTCV 115

Db 63 -----KPDQDITRLVCYNNKITASYEDRVTLPTGITKFSVTR--DTGTYTCWV 111

QY 116 TWQTPDGNQVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCCAR--GSPPI 174

Db 112 SEEGNSYGEYKVKLLIYL-----VPPSKPTVNIPS-----SATIGNRAVLTCSEQDGSPPSE 163

QY 175 YIWKV---QQTN-----NQEPIKVAITLLEKPAVIADSGSVFCTAKGVGSSQH 222

Db 164 YTWFKDGIIVMTPNPKTRAFNSSYVNLPTTGGELVFDPFLSASDTGEYSCEARNGYGPMT 223

QY 223 SDIVK 227

Db 224 SNAVR 228

RESULT 2

138346

elastic titin - human (fragment)
C:Species: Homo sapiens (man)
C:Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 09-Jul-2004
C:Accession: I38346
R:Labait, S.; Kolmerer, B.
Science 270, 293-296, 1995
A:Title: Titins: giant proteins in charge of muscle ultrastructure and elasticity.
A:Reference number: A57430; MUID:96026330; PMID:7569978
A:Accession: I38346
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: mRNA
A:Residues: 1-7962 <RES>
A:CROSS-references: UNIPROT:Q10465; EMBL:X90569; NID:g1017426; PIDN:CAA62189.1; PID:g1017426
C:Genetics:
A:Gene: GDB:TTN
A:CROSS-references: GDB:127867; OMIM:188840
A:Map position: 2q31-2q31

Query Match 10.2%; Score 172; DB 2; Length 7962;
Best Local Similarity 27.5%; Pred. No. 0.00045;
Matches 67; Conservative 30; Mismatches 85; Indels 62; Gaps 11;

QY 16 DTYGRPILEVPES-VTGPWKGDVNLFP-----CTYDPLQGYTVQLVKW-----LVQRGSDP 64
DB 860 DSSGALIVPEPSPFVTKPGSKDV-LPGSAVCLKSTFGSTPLTIRWFKGNKELVSGGSCY 918
QY 65 VTIFLRDSSGDHIQAKYQGRUHLVSHKVPGDVSLQLSTLEMDRSHYTCVETWQTPDG-- 122
DB 919 IT-----KEALESLELYLVKTSDSGTYTCKVS-NVAGGVE 953
QY 123 ---NQVVRDKITELRVOKLSVSKPTVTGSGYGFVTPQGMRLSLOCAQSGSPISYIWK 179
DB 954 CSANLFEVKEPAT--FVEKLEPSQ-----LLKGDATQLACKVTGTPPIKITWFA 1000
QY 180 QQTNNQEPK-----VATLSTLLFKPAVIADSGSYFCTAKGQVSGEHSIDIVKVFVKDSS 234
DB 1001 NDREIKESSKRWMSFVESTAVLRLTDVGIEDSGEYMCEAQAEGSDHCSIV--IVKESP 1058
QY 235 KLLK 238
DB 1059 YFTK 1062

RESULT 3
JC7780
cox sackie- and adenovirus receptor - bovine
C:Species: Bos primigenius taurus (cattle)
C:Date: 02-Apr-2002 #sequence_revision 02-Apr-2002 #text_change 09-Jul-2004
C:Accession: JC7780
R:Theoelen, I.; Keyaerts, E.; Lindberg, M.; Van Ranst, M.
Biochem. Biophys. Res. Commun. 288, 805-808, 2001
A:Title: Characterization of a cDNA encoding the bovine coxsackie and adenovirus receptor
A:Reference number: JC7780
A:Contents: Liver
A:Accession: JC7780
A:Molecule type: mRNA
A:Residues: 1-365 <IHO>
A:CROSS-references: UNIPROT:Q8WMV3; GB:AY033651
C:Comment: This protein serves as the primary adenoviral attachment site on bovine cells

Query Match 10.0%; Score 169; DB 2; Length 365;
Best Local Similarity 24.8%; Pred. No. 1.6e-05;
Matches 85; Conservative 40; Mismatches 120; Indels 98; Gaps 17;

QY 1 MGIILGLLLGLHLTVDYGRPILEVPESVTGPWKGD-VNLPCY-----DPLQGYTVQ 52
DB 1 MELLRLFLLLCGVADFTRGLSI-TTPEQMIERAKGETAYLPCKFTLGPEDQGPLD----- 54
QY 53 LVKVLVQSGS-----DPVTIFLRDSSGDHIQAKYQ---GRHLVSHK--VPGDVSLQLSTL 103
DB 55 -IEWLLSPADNQKVQDVIILY---SGDKIYDDYQDLKGRVHFTSNDLKGSDASINVNL 110
QY 104 EMDRSHYTCEVTWQTPDGNQVVRDKITELRVOKLSVSKPTVTGSGYGFVTPQ-----GM 159

DB 111 QLSDIGTYQCKVKAPGVGNK-----XIQTLTVLKP-----SGIRCYVDGSEELGN 156
QY 160 RISLQOQAR-GSPPISYIWKYQQTNNQEP-----IKVATLSTLLFKPAVIADSGSYFCTAKG 215
DB 157 DFKLKCEPKESGLPLRYEWQKLSDSQKLPTSWLPEMTSPVISVKNASAEYSGTYTCTVRN 216
QY 216 QVSGEQHSDIVKVVVQSSKLLTKTKTEAPTMTYPLKATSVTKQSDWTTTMDMDGVLGTS 275
DB 217 RVGSDQ-----CLLRDLVVPPSNR-----AGTI 239
QY 276 AGPGKSLPVFAIILIISLCMVFTMAXIMLCRKTSGQEHVYE 318
DB 240 AG-----AVIGTLALVLIALIVF-----CCHKGRREEKYE 270

RESULT 4
B49120
protein-tyrosine kinase (EC 2.7.1.112) dtk2 - fruit fly (Drosophila melanogaster)
C:Species: Drosophila melanogaster
C:Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 16-Aug-2004
C:Accession: B49120; S18010
R:Shishido, E.; Higashijima, S.; Emori, Y.; Saigo, K.
Development 117, 751-761, 1993
A:Title: Two FGF-receptor homologues of Drosophila: one is expressed in mesodermal primordium and the other in the nervous system.
A:Reference number: A49120; MUID:93321617; PMID:8330538
A:Accession: B49120
A:Status: preliminary
A:Molecule type: nucleic acid
A:Residues: 1-1052 <SHI>
A:CROSS-references: UNIPROT:Q09147; GB:X74031; GB:S63797; NID:g397600; PIDN:CAA52190.1;
A:Experimental source: pupa
A:Note: sequence extracted from NCBI backbone (NCBIN:135151, NCBIP:135153)
R:Shishido, E.; Emori, Y.; Saigo, K.
FEBS Lett. 289, 235-238, 1991
A:Title: Identification of seven novel protein-tyrosine kinase genes of Drosophila by the complementary DNA method.
A:Reference number: S17552; MUID:92008631; PMID:1915852
A:Accession: S18010
A:Status: not compared with conceptual translation
A:Molecule type: DNA
A:Residues: 869-922 <SH2>
C:Genetics:
A:Gene: FlyBase:bt1; dtk2
A:CROSS-references: FlyBase:FBgn0005592
C:Superfamily: protein kinase homology
C:Keywords: ATP; growth factor receptor; phosphotransferase; tyrosine-specific protein kinase
F:710-1003/Domain: protein kinase homology <KIN>
F:718-726/Region: protein kinase ATP-binding motif

Query Match 8.5%; Score 143.5; DB 2; Length 1052;
Best Local Similarity 23.4%; Pred. No. 0.0055;
Matches 71; Conservative 44; Mismatches 119; Indels 69; Gaps 14;

QY 16 DTYGRPILE-----VPESVTGPWKGDVNLPCY-YDP-LQGYTVQLVKVLVQRGSDP 64
DB 130 DLFPQLNESRLKLLQPLPKTVQRTAGLQQLNCSPNDPAKG---VNISWL----- 179
QY 65 VTIFLRDSSGDHIQAKYQGRHLVSHKVPGDVSLQLSTLEMDRSHYTCVETWQTPDGNQ 124
DB 180 -----HIDTQILGGRGRKIKL---RWSLTVGLOQPEDAGSYHCELCVE---QD 221
QY 125 VVRDKITELRVOKLSVSKPTVTGSGYGFVTPQGMRLSLOCAQSGS---PPISYIWKYQ 181
DB 222 CORSNPTQLEVISRKHVTPMLKPGFPRNTSIALGDNVSIACLEDSALEPKITWL-HKGN 280
QY 182 TNN-----QEPKIVATLSTLTFKPAVI-----ADSGSYFCTAKGQVSGSQH 222
DB 281 ADNIDDLLQRLREQSLPVDVTRLITRMDEPQVLRGLGNVLMEDGGWYICIAENQVGRVA 340
QY 223 SDIVKVFVKDSSKLLTKTKTEAPTMTYPLKATSVTKQSDWTTTMDMDGVLGTSAGPKSL 282
DB 341 ASYVDLYSPSDDTTTTRITTT--TTTVASPIPTASTGEONDD---DVENPAADASGVGVP-- 393

QY 283 PVF 285
DB 394 PVF 396

RESULT 5
S18252
heparan sulfate proteoglycan - mouse
N:Alternate names: perlecan
C:Species: Mus musculus (house mouse)
C:Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 12-Jul-2004
C:Accession: S18252; A31917; B31917; S66460
R:Noonan, D.M.; Fulle, A.; Valence, P.; Cal, S.; Horigan, E.; Sasaki, M.; Yamada, Y.; Hase
J. Biol. Chem. 266, 22939-22947, 1991
A:Title: The complete sequence of perlecan, a basement membrane heparan sulfate proteogly
adhesion molecule.
A:Reference number: S18252; MUID:92078153; PMID:1744087
A:Accession: S18252
A:Molecule type: mRNA
A:Residues: 1-3707 <NOO>
A:Cross-references: UNIPROT:Q05793; EMBL:M77174; NID:g200295; PIDN:AAA39911.1; PID:g2002
R:Noonan, D.M.; Horigan, E.A.; Ledbetter, S.R.; Vogeli, G.; Sasaki, M.; Yamada, Y.; Hase
J. Biol. Chem. 263, 16379-16387, 1988
A:Title: Identification of cDNA clones encoding different domains of the basement membra
A:Reference number: A92680; MUID:89034110; PMID:2972708
A:Accession: A31917
A:Molecule type: mRNA
A:Residues: 940-1601 <NO2>
A:Cross-references: GB:J04054; NID:g200252; PIDN:AAA39899.1; PID:g200253
A:Accession: B31917
A:Molecule type: protein
A:Residues: 1272-1274, 'X', 1276, 'X', 1278-1279 <SCH>
C:Keywords: glycoprotein
F:199-234/Domain: LDL receptor ligand-binding repeat homology <LDL1>
F:285-319/Domain: LDL receptor ligand-binding repeat homology <LDL2>
F:325-359/Domain: LDL receptor ligand-binding repeat homology <LDL3>
F:368-403/Domain: LDL receptor ligand-binding repeat homology <LDL4>
F:764-811/Domain: laminin-type EGF-like homology <LEG>
F:1159-1206/Domain: laminin-type EGF-like homology <LEG7>
F:1563-1610/Domain: laminin-type EGF-like homology <LEG8>
F:1613-1668/Domain: laminin-type EGF-like homology <LEG9>
F:3163-3198/Domain: EGF homology <EGF>
F:3270-3423/Domain: laminin G repeat homology <LG2>
F:3464-3492/Domain: EGF homology <EGF7>
F:1256, 1891, 2336, 2394, 2427/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 8.3%; Score 139.5; DB 2; Length 3707;
Best Local Similarity 25.1%; Pred. No. 0.055;
Matches 66; Conservative 31; Mismatches 105; Indels 61; Gaps 14;

QY 13 LTVDT-----YGRPILVEPESVTPGWKGVNLPCTYDPLQGYTVLVKLVQSGDPVITF 68
DB 2521 LTVDTGVAPGPPQVVEES-----ELTLEA-----GHTATL--HCSATGNPPPT-- 2563

QY 69 LRDSGDHIQAKYQGRHSHVSKVPGDVSLQSLTLEMDRSHYTCVWTQPDGQNVVD 128
DB 2564 -----IHWKLRAPLPWQRIEGLN-TLVIPRAQDSQGYICNAT-----NSAGHT 2608

QY 129 KTELVRVQKLSVKPTVTGSGYGTVPQ-----GMRISLQCGARGSPPIYSIYWKQOT 182
DB 2609 EATV-----LHVESPP-----YATIIPEHTSAQPNLVQLCLAHGTPPLTYQW--SLV 2656

QY 183 NNQEPKIVATLSTIL-FKPAVIADSGSYFCTAKGVGSEQSHSDIVKVVQKSSKLLKTKT 241
DB 2657 GGVLPKAVVRNQLLRLEPTPEDSGRYRCQVSNRVGSAE--AFAQVILVQSSSNLPDTS 2714

QY 242 ----EAPTTMTYPLKATSTVKOS 260
DB 2715 IPGSGTPTVQVTPQLETRNIGAS 2737

RESULT 6
PN0568
connectin 3B - chicken (fragment)
N:Alternate names: Cn3B protein
C:Species: Gallus gallus (chicken)
C:Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 09-Jul-2004
C:Accession: PN0568
R:Maruyama, K.; Endo, T.; Kume, H.; Kawamura, Y.; Kanzawa, N.; Nakauchi, Y.; Kimura, S.;
Biochem. Biophys. Res. Commun. 194, 1288-1291, 1993
A:Title: A novel domain sequence of connectin localized at the 1 band of skeletal muscle
A:Reference number: PN0568; MUID:93356802; PMID:8352787
A:Accession: PN0568
A:Molecule type: mRNA
A:Residues: 1-1323 <MAR>
A:Cross-references: UNIPROT:Q08476; DBJ:D16541; NID:g391629; PID:d1004495; PID:g391630
A:Experimental source: skeletal muscle
C:Comment: This protein string-like single molecule spans from the Z line to the M line

Query Match 8.2%; Score 138.5; DB 2; Length 1323;
Best Local Similarity 21.4%; Pred. No. 0.018;
Matches 61; Conservative 38; Mismatches 117; Indels 69; Gaps 10;

QY 22 ILEVPESV---TGPWKGVNLPCTYDPLQGYTVLVKLVQSGDPVITFLRDSGDHIQ 78
DB 143 ILEIPNSKLEDOGGYQCHIENDSGDNCHGAITILEPPYFVTPLPEPVQTVGDSASLQ 202

QY 79 QAK-----YQG-----RLVSHKVPGVDSVLQSLTLEMDRSHYTCVWTQTP 120
DB 203 VAGTPEMIVSYKGDGTLRGATATVMHFNQV---ATLVFQVSDSDSGEVICKVENTVG 259

QY 121 DGNQ-----VVRDKITELRVQKLSVKPTVTGSGYGTVPQGMRIISLQCGARGSPPIYS 175
DB 260 EATSSLLTVQERKLPPSFRKLRDVHETV-----GLPVTDCGIAGEPIEV 307

QY 176 IWKYQNTNNQEPKIVAT--LSTLFPKPAVIADS---GSYFCTAKGVGSEQSHSDIVKFW 230
DB 308 SWFKDNVRVKEDYVHTSFIDNVAAILQLTKSLMGQYTCAGNAIGT-----A 357

QY 231 KDSKLLKTKTEAPTTMTYPLKATSTVKQSWDVTMDGVLGETS 275
DB 358 SSSGKLVLTGKTPPPFDTPI-----TPVDGIIGESA 389

RESULT 7
T42633
connectin/titin - chicken (fragment)
C:Species: Gallus gallus (chicken)
C:Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 09-Jul-2004
C:Accession: T42633
R:Yajima, H.; Ohtsuka, H.; Kawamura, Y.; Kume, H.; Murayama, T.; Abe, H.; Kimura, S.; Ma
Biochem. Biophys. Res. Commun. 223, 160-164, 1996
A:Title: A 11.5-kb 5'-terminal cDNA sequence of chicken breast muscle connectin/titin re
A:Reference number: 222221; MUID:96254045; PMID:8660363
A:Accession: T42633
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: mRNA
A:Residues: 1-4162 <YAJ>
A:Cross-references: UNIPROT:Q98918; EMBL:D83390; NID:gi513029; PIDN:BAAL1908.1; PID:gi51
A:Experimental source: breast muscle
C:Keywords: skeletal muscle

Query Match 8.2%; Score 138.5; DB 2; Length 4162;
Best Local Similarity 21.4%; Pred. No. 0.076;
Matches 61; Conservative 38; Mismatches 117; Indels 69; Gaps 10;

QY 22 ILEVPESV---TGPWKGVNLPCTYDPLQGYTVLVKLVQSGDPVITFLRDSGDHIQ 78
DB 2715 IPGSGTPTVQVTPQLETRNIGAS 2737

Db 3895 ILEIPNSKLEDOGOYSCHIEENDSGQDNCHGAITILEPPYFVTPLEPQVTVGDSASLQCC 3954

Qy 79 QAK-----YQG-----RLVSHKVPDVSLSQLSTLEMDRSHYTCVETWQTP 120

Db 3955 VAGTPEMIVSWYKGTDLRGATVKKHFNQV---ATLVFSQVSDSDSGSEYICKVENTVG 4011

Qy 121 DGNQ-----VVRDKITELRVOKLSVSKPTVTGSGYGFVPGQMRISLQCCQARGSPPIISY 175

Db 4012 EATSSLLTVQERKLPPSPFTRKLRDVHEV-----GLPVTDCGIAGSEPIEV 4059

Qy 176 IWYKQQTNNQEPKIVAT--LSTLLFKPAVIADS---GSYFCTAKQGVGSEQHSDIVKFFV 230

Db 4060 SFKDNVRVKEDYNVHTSFIDNVAIQLIKTDKSLMGVYCTCTASNAIGT-----A 4109

Qy 231 KDSSKLLKTKTAPATTMYPLKATSTVKSQSWDWTMDGILGETS 275

Db 4110 SSSGKLVLTGKTPPPFPDTPPI-----TPVDGIIGESA 4141

RESULT 8

T08678

hypothetical protein DKF2p564I11922.1 - human (fragment)

C:Species: Homo sapiens (man)

C>Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004

C:Accession: T08678

R:Wambutt, R.; Heubner, D.; Mewes, H.W.; Gassenhuber, J.; Wiemann, S.

submitted to the Protein Sequence Database, May 1999

A:Reference number: Z16469

A:Accession: T08678

A:Molecule type: mRNA

A:Residues: 1-584 <WAM>

A:Cross-references: UNIPROT:Q9Y3Y8; EMBL:AL049946

A:Experimental source: fetal brain; clone DKF2p564I11922

C:Genetics:

A:Note: DKF2p564I11922.1

Query Match 8.1%; Score 136; DB 2; Length 584;

Best Local Similarity 22.4%; Pred. No. 0.0099;

Matches 53; Conservative 37; Mismatches 99; Indels 48; Gaps 8;

Qy 35 GDNLPCTYDPIQGYTVQLVKLVQRGSDPVTIFLRDSSGDHIQQAQYQGRLHVSHKVP 94

Db 15 GOLKVDVCV---ATGLPNPEISMSLPDGSIVNSFMQSDSDSGRTKR-----YVFN 61

Qy 95 DVSLSQLSTLEMDRSHYTCVETWQTPDGNQVVRDKITELRVOKLSVSKP-TVTTGSGYGF 153

Db 62 NGTLFNEVGMREEGDYTCFAE-----NQVGKD---EMVRVKVVTAPATIRNKTCLAV 112

Qy 154 TVPQGMRLISLQCCARGSPPIISYIWY---KQQTNNQEPKIVATLSTLLFKPAVIADSGSY 209

Db 113 QVPYGDVTVVACEAKGEMPKVTWLSPTNKV IPTSSEKVIQIYQDGTLLIQKQRSDSGNY 172

Qy 210 FCTAKQGVGSEQH-----SDIVKPVVXKSSKLLKTKTEA-PT 245

Db 173 TCLVRNSAGEDRKTWIIHVNVQPPKINGPNPITTVREIAAGGSKLIDCKAEGIPT 229

RESULT 9

T20992

hypothetical protein F15G9.4a - Caenorhabditis elegans

C:Species: Caenorhabditis elegans

C>Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004

C:Accession: T20992; T24733

R:Sulston, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19355

A:Accession: T20992

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5175 <WIL>

A:Cross-references: UNIPROT:Q8I0J3; EMBL:Z47068; PIDN:CAA87335.1; GSPDB:GN000028; CESP:F1

A:Experimental source: clone F15G9

R:Kershaw, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19929

A:Accession: T24733

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5175 <WI2>

A:Cross-references: EMBL:Z47070; PIDN:CAA87344.1; GSPDB:GN000028; CESP:F15G9.4a

A:Experimental source: clone T09B9

C:Genetics:

A:Gene: CESP:F15G9.4a

A:Map position: X

A:Introns: 85/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184/3; 120/1; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 5077/1;

Query Match 7.9%; Score 133.5; DB 2; Length 5175;

Best Local Similarity 22.9%; Pred. No. 0.24;

Matches 48; Conservative 36; Mismatches 93; Indels 33; Gaps 6;

Qy 76 HIQQAQYQGRLHVSHKVPDVSLSQLSTLEMDRSHYTCVETWQTPDGNQVVRDKITELRV 135

Db 2796 HAHDESQVNGVITSKYAAANEKTLNVTNIQLDDEGFYIC-----TAVNEAGITKKFFKLIV 2850

Qy 136 -----QKLSVSKPTVTGSGYGFVPGQMRISLQCCQARGSPPIISYIWYKQQTNNQE 186

Db 2851 IETPYFLDQQL-----YPIILGKRLTLDCSATGTPPTILFMKDGKRLNE 2896

Qy 187 PIKVATL-STLLFKPAVIADSGSYFCTAKQGVGSEQHSDIVKFFV--KDSKLLKTKTEA 243

Db 2897 SDEVDIIGSTLVIDNPQKEVEGRYTCTIAENKAGRSEKDMVMVEVLLPPLKSKWINVEVQA 2956

Qy 244 --PTTMYPLKATSTVKSQSWDWTMDGYL 271

Db 2957 GDPLTLECPIDETSGVHITWSRQFGKDGQL 2986

RESULT 10

T43290

hemiceitin precursor - Caenorhabditis elegans

C:Species: Caenorhabditis elegans

C>Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 09-Jul-2004

C:Accession: T43290; T20993; T24734

R:Voegel, B.E.; Hedgecock, E.M.

submitted to the EMBL Data Library, June 1998

A:Description: Hemiceitin is required for hemidesmosome mediated cell adhesion and germ-

A:Reference number: Z22396

A:Accession: T43290

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: mRNA

A:Residues: 1-5198 <VOG>

A:Cross-references: UNIPROT:O76518; EMBL:AF074901; PIDN:AAC26792.1

R:Sulston, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19355

A:Accession: T20993

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5198 <WIL>

A:Cross-references: EMBL:Z47068; PIDN:CAA87336.1; GSPDB:GN000028; CESP:F15G9.4b

A:Experimental source: clone F15G9

R:Kershaw, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19929

A:Accession: T24734

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5198 <WI2>

A:Cross-references: EMBL:Z47070; PIDN:CAA87345.1; GSPDB:GN000028; CESP:F15G9.4b

A:Experimental source: clone T09B9

C:Genetics:

A:Gene: him-4; F15G9.4b

A:Map position: X

A:Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184/3;

2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/1
1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4902/1; 4964/1; 5034/1; 5100/

Query Match 7.9%; Score 133.5; DB 2; Length 5198;
Best Local Similarity 22.9%; Pred. No. 0.24;
Matches 48; Conservative 36; Mismatches 93; Indels 33; Gaps 6;

QY 76 HQQAQYQGRHLVSHKVPDVSLSQLEMDRSHYTCVTTWTPDGNQVVRDKITELRV 135
DB HAHDSEVQNGVITSKYAANEKTLNVTNIQLDDSGFYFC-----TAVNEAGITKKFKKLIV 2850

QY 136 -----QKLSVSKPTVTTCGGYGTVPQGNRISLQCAQSGSPISVIWYKQNTNOE 186
DB IETPYFLDQOK-----YPIILGKRLTDCSATGTPPTILFMKQGRKLN 2896

QY 187 PIKVA TL-STLFPKPAVIADSGSYFTAKQGVGEHQSDIVKPVV--KDSSKLLTKTKTEA 243
DB SDEVDIIGSTLVIDNQKEVGRYTCIAENKAGSEKMMVEVLLPPLSKSEWINVEVQA 2956

QY 244 --PTTWYPLKATSVKQSWDWTMDGYL 271
DB GDPLTLECIPTDTSVGHITMSRQFGKDGQL 2986

RESULT 11

PFHUGB
N: Contains: protein-tyrosine kinase (EC 2.7.1.112)
C: Species: Homo sapiens (man)
C: Date: 31-Dec-1992 #sequence_revision 31-Dec-1992 #text_change 09-Jul-2004
C: Accession: A8206; A31195; A38268; B31925; C31925
R: Gronwald, R.G.K.; Grant, F.J.; Haldeman, B.A.; Hart, C.E.; O'Hara, P.J.; Hagen, F.S.;
Proc. Natl. Acad. Sci. U.S.A. 85, 3435-3439, 1988
A: Title: Cloning and expression of a cDNA coding for the human platelet-derived growth factor
A: Reference number: A8206; MUID: 88217915; PMID: 2835772
A: Accession: A8206
A: Molecule type: mRNA
A: Residues: 1-1106 <GRO>
A: Cross-references: UNIPROT: P09619; GB: J03278; NID: g189731; PIDN: AAA60049.1; PID: g189732
R: Claesson-Welsh, L.; Eriksson, A.; Moren, A.; Severinsson, L.; Ek, B.; Oestman, A.; Bet
Mol. Cell. Biol. 8, 3476-3486, 1988
A: Title: cDNA cloning and expression of a human platelet-derived growth factor (PDGF) re
A: Reference number: A31195; MUID: 89096941; PMID: 2850496
A: Accession: A31195
A: Molecule type: mRNA
A: Residues: 1-240.'D'. 242-1106 <CLA>
A: Cross-references: GB: M21616; NID: g189729; PIDN: AAA36427.1; PID: g189730
R: Partanen, J.; Mäkelä, T.P.; Alitalo, R.; Lehtvaesaiho, H.; Alitalo, K.
Proc. Natl. Acad. Sci. U.S.A. 87, 8913-8917, 1990
A: Title: Putative tyrosine kinases expressed in K-562 human leukemia cells.
A: Reference number: A38268; MUID: 91062389; PMID: 2247464
A: Accession: A38268
A: Status: nucleic acid sequence not shown
A: Molecule type: mRNA
A: Residues: 828-884 <PAR>
R: Roberts, W.M.; Look, A.T.; Roussel, M.F.; Sherr, C.J.
Cell 55, 655-661, 1988
A: Title: Tandem linkage of human CSF-1 receptor (c-fms) and PDGF receptor genes.
A: Reference number: A9098; MUID: 89028677; PMID: 2846185
A: Accession: A31925
A: Status: not compared with conceptual translation
A: Molecule type: DNA
A: Residues: 676-727 <ROB>
A: Accession: B31925
A: Status: not compared with conceptual translation
A: Molecule type: DNA
A: Residues: 901-932 <R02>
A: Accession: C31925
A: Status: not compared with conceptual translation
A: Molecule type: DNA
A: Residues: 1047-1106 <R03>
C: Comment: The extracellular domain is predicted to include five immunoglobulin-like dom
C: Genetatics:

A: Gene: GDB: PDGFRB
A: Cross-references: GDB: 120710; OMIM: 173410
A: Map position: 5q31-5q32
C: Superfamily: macrophage colony-stimulating factor 1 receptor; immunoglobulin homology,
C: Keywords: ATP; autophosphorylation; glycoprotein; growth factor receptor; heterodimer,
F1-32/Domain: signal sequence #status predicted <SIG>
F33-1106/Product: platelet-derived growth factor receptor beta #status predicted <EXT>
F33-531/Domain: extracellular #status predicted <IMM1>
F47-102/Domain: immunoglobulin homology <IMM2>
F142-192/Domain: immunoglobulin homology <IMM2>
F228-293/Domain: immunoglobulin homology <IMM3>
F429-510/Domain: immunoglobulin homology <IMM4>
F532-555/Domain: transmembrane #status predicted <TM>
F556-1106/Domain: intracellular #status predicted <INT>
F598-965/Domain: protein kinase homology <KIN>
F606-614/Region: protein kinase ATP-binding motif
F45-89, 103, 215, 230, 292, 307, 354, 371, 468, 479/Binding site: carbohydrate (Aan) (covalent)
F54-100, 149-190, 235-291, 436-508/Disulfide bonds: #status predicted
F:54/Active site: Lys #status predicted
F:837/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

Query Match 7.9%; Score 133; DB 1; Length 1106;
Best Local Similarity 21.4%; Pred. No. 0.038;
Matches 60; Conservative 41; Mismatches 92; Indels 88; Gaps 11;

QY 23 LEVPESVTGPW-----KGDVNLPCYDPLQYQTVLVKLVQRGSDPVTIFLRDSSG 74
DB ITIPCRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 75 DHQQAQYQGRHLVSHKVPDVSLSQLEMDRSHYTCVTTWTPDGNQVVRDKITELR 134
DB -----FEDRSYIKTTIGD-----REVSDDAY-----VYRLQ 210

QY 135 VQKLSVSKPTVTTCGGYGTVPQGNRISLQCAQSGSPISYIW-YKQNTNO--EPIKVA 191
DB VSSINVSNAVQT-----VVRQGENITLMCIIVGNEVNVFETPRKESGRLEVPVTF 264

QY 192 TL-----STLLFPKPAVIADSGSYFTAKQGVGEHQSDIVKPVVYKDSKLLTKTKTEAP 244
DB LLDMPVHRSILHPSAELESDGTYTCNVTESVNDHQDEKAINITVBSG-VYRLGEGV 323

QY 245 TWTVPKATSVKQSWD-----WTDMDGYLGETSAG 277
DB TLQFAELHRSRTLQVVFAYPPPTVLWFKD-NRTLGDSSAG 363

RESULT 12

C42632
cell adhesion molecule apCAM (clone dl2) - California sea hare
C: Species: Aplysia californica (California sea hare)
C: Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C: Accession: C42632
R: Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
Science 256, 638-644, 1992
A: Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plasti
A: Reference number: A42632; MUID: 92263095; PMID: 1585176
A: Accession: C42632
A: Status: preliminary; not compared with conceptual translation
A: Molecule type: nucleic acid
A: Residues: 1-765 <NAY>
A: Cross-references: UNIPROT: Q9TWA4
A: Experimental source: CNS
A: Note: sequence extracted from NCBI backbone (NCBI:101351)
C: Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; imm

Query Match 7.8%; Score 132.5; DB 2; Length 765;
Best Local Similarity 20.7%; Pred. No. 0.026;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHYTCV-----TWQTPDGNQVVRDKITELRQKLSVSKPT-----VTTGSG 150
DB KVGDEVKITCOATGVPPPTYPQFKGDVWVTDENVNGVLTINPLKTTDQATTCIATNKG 289

QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
Db 290 -GFAESSNTLDVKVPPTIEDMEETYDAVSGQELTITCTAKGDPPEPSVIWKDGPQASSTD 348
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSDIVKF 228
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
QY 229 VVK-----VP-----DSSKLLKTKT 241
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWYMPDAENPDDYSKAVRIEN 466
QY 242 EAPTMTYPLKATSTVKQSWD 262
Db 467 EAPYTI-----NMLQKWD 479

RESULT 13
B42632
C:Species: Aplysia californica (clone d15) - California sea hare
C:Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C:Accession: B42632
R:Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
S:Science 256, 638-644, 1992
A:Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plasticity
A:Reference number: A42632; MUID:92263095; PMID:1585176
A:Accession: B42632
A>Status: preliminary; not compared with conceptual translation
A:Molecule type: nucleic acid
A:Residues: 1-812 <MAY>
A:Cross-references: UNIPROT:Q9TWA5
A:Experimental source: CNS
A>Note: sequence extracted from NCBI backbone (NCBIP:101346)
C:Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; immunoglobulin domain

Query Match 7.8%; Score 132.5; DB 2; Length 812;
Best Local Similarity 20.7%; Pred. No. 0.028;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHTYCEV-----TWQTPDGNQVVRDKITELRVQKLSVSKPT-----VTGSG 150
Db 230 KVGDEVKITCQATGVPPPTYQFKGDMVTDEMVGNGVLTINPLKTTDQATVTCIATNKG 289
QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
Db 290 -GFAESSNTLDVKVPPTIEDMEETYDAVSGQELTITCTAKGDPPEPSVIWKDGPQASSTD 348
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSDIVKF 228
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
QY 229 VVK-----VP-----DSSKLLKTKT 241
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWYMPDAENPDDYSKAVRIEN 466
QY 242 EAPTMTYPLKATSTVKQSWD 262
Db 467 EAPYTI-----NMLQKWD 479

RESULT 14
A42632
C:Species: Aplysia californica (clone d19) - California sea hare
C:Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C:Accession: A42632
R:Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
S:Science 256, 638-644, 1992
A:Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plasticity
A:Reference number: A42632; MUID:92263095; PMID:1585176
A:Accession: A42632
A>Status: preliminary; not compared with conceptual translation

A:Molecule type: nucleic acid
A:Residues: 1-932 <MAY>
A:Cross-references: UNIPROT:Q9TWA6
A:Experimental source: CNS
C:Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; immunoglobulin domain

Query Match 7.8%; Score 132.5; DB 2; Length 932;
Best Local Similarity 20.7%; Pred. No. 0.033;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHTYCEV-----TWQTPDGNQVVRDKITELRVQKLSVSKPT-----VTGSG 150
Db 230 KVGDEVKITCQATGVPPPTYQFKGDMVTDEMVGNGVLTINPLKTTDQATVTCIATNKG 289
QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
Db 290 -GFAESSNTLDVKVPPTIEDMEETYDAVSGQELTITCTAKGDPPEPSVIWKDGPQASSTD 348
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSDIVKF 228
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
QY 229 VVK-----VP-----DSSKLLKTKT 241
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWYMPDAENPDDYSKAVRIEN 466
QY 242 EAPTMTYPLKATSTVKQSWD 262
Db 467 EAPYTI-----NMLQKWD 479

RESULT 15
S26180
C:Species: Gallus gallus (chicken)
C:Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C:Accession: S26180
R:Volkmmer, H.; Hassel, B.; Wolff, J.M.; Frank, R.; Rathjen, F.G.
J. Cell Biol. 118, 149-161, 1992
A:Title: Structure of the axonal surface recognition molecule neurofascin and its relationship to the cell adhesion molecule L1
A:Reference number: S26180; MUID:92317154; PMID:1377696
A:Accession: S26180
A>Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-1272 <VOL>
A:Cross-references: UNIPROT:O42414; EMBL:X65224; NID:G63659; PIDN:CAA46330.1; PID:G63660
C:Superfamily: neural cell adhesion molecule L1; fibronectin type III repeat homology; immunoglobulin domain

Query Match 7.8%; Score 132.5; DB 2; Length 1272;
Best Local Similarity 22.2%; Pred. No. 0.049;
Matches 51; Conservative 38; Mismatches 88; Indels 53; Gaps 7;

QY 22 ILEVPESVTPGPKGDVNLPCYDPLQGYTVLKVWLQVORGSQVPTIFLRLDSSGDHITQQA 81
Db 150 VIEVDEG-----APLSLQCNPPP-----GLPPPVIFWSSNEPIHQDK 188
QY 82 -----YQGRHSHVSHKVPDVSLSLS-----TLEMDRSHYTCBVTWTPDGNQVVRDK 129
Db 189 RVSGONGDLYFSNVLQDAQTDYSCNARFHTHTTQQKNPYTLKVKTKKPHNETSLRNH 248
QY 130 ITELVRQKLSVSKPT--VTGSGYGYFTVPOQNRISLQCOARGSPPISYIW----- 179
Db 249 TDMYSGRGVTEPTPFGMYPGYSSQVMVLRGVLDLLECIASGVPAFDINWYKKGELPAG 308
QY 180 --QOTNNOEPIKVATLSTLLFKPAVIADSGSYFCTAKGQVGSQHSDIVK 227
Db 309 KTKLENFNKALRISNVSE-----EDSGEYFCLASNKMGSGIRHTISVR 350

Search completed: August 26, 2005, 19:39:58
Job time : 72 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: August 26, 2005, 19:27:17 ; Search time 170 Seconds

(without alignments)

966.926 Million cell updates/sec

Title: US-10-767-374-2

Perfect score: 1688

Sequence: 1 MGILLGLLLGLHVTDTYGR.....AYIMLCRKTSQOEHYEAAR 321

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1612378 seqs, 512079187 residues

Total number of hits satisfying chosen parameters: 1612378

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 1500 summaries

Database : UniProt_03.*

1: uniprot_spot.*

2: uniprot_trembl.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query %	Match	Length	DB	ID	Description
1	1688	100.0	321	2	Q6UXI4		Q6uxi4 homo sapien
2	1688	100.0	399	2	Q9Y279		Q9y279 homo sapien
3	728	43.1	280	2	Q8OWA3		Q8owa3 mus musculu
4	178.5	10.6	299	1	JAM1_HUMAN		Q9y624 homo sapien
5	177	10.5	319	1	A33_HUMAN		Q95795 homo sapien
6	172	10.2	7962	2	Q10465		Q10465 homo sapien
7	172	10.2	34350	2	Q8WZ42		Q8wz42 homo sapien
8	170.5	10.1	292	2	Q66I72		Q66i72 brachydanio
9	169.5	10.0	298	1	JAM1_BOVIN		Q9xt56 bos taurus
10	169	10.0	365	2	Q8WMV3		Q8wmv3 bos taurus
11	168	10.0	300	2	Q8VC39		Q8vc39 mus musculu
12	167	9.9	300	1	JAM1_MOUSE		Q88792 mus musculu
13	167	9.9	300	2	Q9UHY1		Q9jhy1 rattus norv
14	167	9.9	319	2	Q9TU79		Q9tu79 sus scrofa
15	166	9.8	319	2	Q8UT80		Q9tu80 canis famil
16	160	9.5	291	2	Q66J15		Q66j15 xenopus tro
17	159.5	9.4	319	1	A33_MOUSE		Q9jka5 mus musculu
18	158	9.4	344	2	Q9UKV4		Q9ukv4 homo sapien
19	158	9.4	365	1	CXAR_HUMAN		P78310 homo sapien
20	157.5	9.3	332	2	Q6P359		Q6p359 xenopus tro
21	156.5	9.3	5516	2	Q7ZZ48		Q7zz48 brachydanio
22	156	9.2	289	2	Q7ZWT0		Q7zwt0 xenopus lae
23	156	9.2	335	2	Q9PWR4		Q9pwr4 gallus gall
24	156	9.2	335	2	Q9YGH1		Q9ygh1 gallus gall
25	153.5	9.1	373	2	Q9H6B4		Q9h6b4 homo sapien
26	153.5	9.1	512	2	Q96DN8		Q96dn8 homo sapien
27	153.5	9.1	5636	2	Q96RW7		Q96rw7 homo sapien
28	151	8.9	390	2	Q9H1X9		Q9h1x9 homo sapien
29	149.5	8.9	344	2	Q9R067		Q9r067 rattus norv
30	149.5	8.9	358	2	Q9R066		Q9r066 rattus norv
31	146.5	8.7	373	2	Q920S5		Q920s5 mus musculu

Q9ygv5 gallus gall
Q8ki90 rattus norv
Q8r373 mus musculu
Q6u7i4 brachydanio
Q7vrf5 canis famil
Q7psj8 anopheles g
Q9vp08 drosophila
Q99147 drosophila
Q8iga5 drosophila
Q90y50 brachydanio
Q99n28 m nectin-li
Q7kvk3 drosophila
Q9w2l3 drosophila
Q44924 drosophila
Q9vuc8 drosophila
Q91w66 mus musculu
P97792 mus musculu
Q9dbj8 mus musculu
Q9xrc3 homo sapien
Q6kdz1 gallus gall
Q7qiv4 anopheles g
Q7tn00 rattus norv
Q05793 mus musculu
Q9vqy0 drosophila
Q9bx67 homo sapien
Q6zn11 homo sapien
Q08476 gallus gall
Q98918 gallus gall
Q804r4 brachydanio
Q9d8b7 mus musculu
Q96896 homo sapien
Q8bu57 mus musculu
Q80ze2 mus musculu
Q920g3 mus musculu
Q96f11 homo sapien
Q967d9 drosophila
Q8iq17 drosophila
Q967d8 drosophila
Q6nr34 drosophila
Q9vqy1 drosophila
Q9vqy2 drosophila
Q967d7 drosophila
Q7kqk5 drosophila
Q9d1m9 mus musculu
Q9epk4 m junctiona
Q24372 drosophila
Q9v6c2 drosophila
Q7qe16 anopheles g
Q8mk6 drosophila
Q8mk7 drosophila
Q9nba1 drosophila
Q8mk8 drosophila
Q9nr99 homo sapien
P97269 cavia porce
Q640u3 xenopus tro
Q63z94 xenopus lae
Q9vt76 drosophila
Q9v3y8 homo sapien
Q9v3y8 drosophila
Q9vgs drosophila
Q8c5k9 mus musculu
Q925f2 mus musculu
Q90ym1 brachydanio
Q8vzh8 rattus norv
Q9erc8 mus musculu
Q8ce95 mus musculu
Q7zxr4 xenopus lae
Q6uxi0 homo sapien
Q6wrh9 rattus norv
Q8av57 gallus gall
Q5wri0 homo sapien
Q6info xenopus lae
Q8ni26 homo sapien
Q9ujp1 homo sapien

105	133.5	7.9	434	2	Q6DN72	Q6dn72 homo sapien	178	127	7.5	1555	2	Q7PPH8	Q7pph8 anopheles g
106	133.5	7.9	484	2	Q25475	Q25475 schistocerc	179	127	7.5	2053	2	Q8XU7	Q8xw7 homo sapien
107	133.5	7.9	1164	2	Q7T2H2	Q7t2h2 gallus gall	180	127	7.5	2092	2	Q76MU9	Q76mu9 homo sapien
108	133.5	7.9	1867	2	Q66MN5	Q66mn5 drosophila	181	127	7.5	2113	2	Q8TD84	Q8td84 homo sapien
109	133.5	7.9	5175	2	Q810L3	Q810l3 caenorhabdi	182	127	7.5	2133	2	Q7PQG9	Q7pqg9 anopheles g
110	133.5	7.9	5198	2	Q76518	Q76518 caenorhabdi	183	126.5	7.5	2133	2	Q6NW88	Q6nw88 brachydanio
111	133	7.9	305	2	Q8VIM2	Q8vim2 mus musculus	184	126.5	7.5	812	2	Q8N612	Q8n612 homo sapien
112	133	7.9	310	2	Q68FQ2	Q68fq2 rattus norv	185	126.5	7.5	865	2	Q68DA2	Q68da2 homo sapien
113	133	7.9	349	1	LACH_SCHAM	Q26474 schistocerc	186	126.5	7.5	924	1	ICAS_HUMAN	Q2umf0 homo sapien
114	133	7.9	394	2	Q7ZX1	Q7zxx1 xenopus lae	187	126.5	7.5	924	2	Q8TAM9	Q8tam9 homo sapien
115	133	7.9	697	2	Q8NC72	Q8nc72 homo sapien	188	126.5	7.5	1056	2	Q90Z03	Q90z03 xenopus lae
116	133	7.9	1059	2	Q6UKL7	Q6uxl7 homo sapien	189	126	7.5	318	2	Q91664	Q91664 xenopus lae
117	133	7.9	1106	1	PGDR_HUMAN	P09619 homo sapien	190	126	7.5	413	2	Q7QBV2	Q7qbv2 anopheles g
118	133	7.9	1119	2	Q6UXM1	Q6uxm1 homo sapien	191	126	7.5	885	2	Q8HYV1	Q8hyv1 sus scrofa
119	132.5	7.8	298	2	Q9J159	Q9j159 m vascular	192	126	7.5	886	2	Q8HYV2	Q8hyv2 sus scrofa
120	132.5	7.8	463	2	Q66J72	Q66j72 xenopus lae	193	126	7.5	948	2	Q9VME2	Q9vme2 drosophila
121	132.5	7.8	765	2	Q9BKQ1	Q9bkq1 aphysia cal	194	126	7.5	1235	2	Q7Q0S7	Q7q0s7 anopheles g
122	132.5	7.8	765	2	Q9TWA4	Q9twa4 aphysia cal	195	126	7.5	1431	2	Q80U60	Q80u60 mus musculus
123	132.5	7.8	812	2	Q9BKQ0	Q9bkq0 aphysia cal	196	125.5	7.4	303	2	Q7Q154	Q7q154 anopheles g
124	132.5	7.8	812	2	Q9TWA5	Q9twa5 aphysia cal	197	125.5	7.4	412	2	Q9R1E1	Q9r1e1 rattus norv
125	132.5	7.8	932	2	Q9BKp9	Q9bkp9 aphysia cal	198	125.5	7.4	707	2	Q7PMJ1	Q7pmj1 anopheles g
126	132.5	7.8	932	2	Q9TWA6	Q9twa6 aphysia cal	199	125.5	7.4	1443	2	Q8MTB2	Q8mtb2 drosophila
127	132.5	7.8	1369	1	NFAS_CHICK	Q42414 gallus gall	200	125.5	7.4	1496	2	Q92626	Q92626 homo sapien
128	132	7.8	300	2	Q640C0	Q640c0 xenopus lae	201	125.5	7.4	1765	2	Q9VS30	Q9vs30 drosophila
129	132	7.8	302	2	Q7SYQ7	Q7syq7 xenopus lae	202	125.5	7.4	1770	2	Q9VS29	Q9vs29 drosophila
130	132	7.8	394	2	Q6AYD4	Q6ayd4 rattus norv	203	125	7.4	313	2	Q57596	Q57596 gallus gall
131	132	7.8	1746	2	Q8WY19	Q8wy19 homo sapien	204	125	7.4	315	2	Q9DGI5	Q9dgi5 gallus gall
132	132	7.8	2012	1	DSCA_HUMAN	Q60469 homo sapien	205	125	7.4	1006	2	Q6ID89	Q6ide9 drosophila
133	131.5	7.8	461	2	Q13854	Q13854 homo sapien	206	124.5	7.4	259	2	Q7Z2Q1	Q7z2q1 homo sapien
134	131.5	7.8	1117	2	Q6P1C6	Q6p1c6 mus musculus	207	124.5	7.4	344	2	Q93242	Q93242 gallus gall
135	131.5	7.8	1946	2	Q68J72	Q68j72 apis mellif	208	124.5	7.4	353	1	CEPU_CHECK	Q96773 gallus gall
136	131.5	7.8	2673	2	Q96SC3	Q96sc3 homo sapien	209	124.5	7.4	387	2	Q86XK7	Q86xk7 homo sapien
137	131	7.8	259	2	Q9Y5B2	Q9y5b2 homo sapien	210	124.5	7.4	412	2	Q8NZS4	Q8nzs4 homo sapien
138	131	7.8	285	2	Q9D780	Q9d780 mus musculus	211	124.5	7.4	877	2	Q9GSH3	Q9gsh3 halocynthia
139	131	7.8	714	2	Q6ZPE6	Q6zpe6 mus musculus	212	124.5	7.4	1073	2	Q9W1T8	Q9wit8 drosophila
140	131	7.8	4391	1	PGBM_HUMAN	P98160 homo sapien	213	124.5	7.4	1173	2	Q6NR54	Q6nr54 drosophila
141	130.5	7.7	272	2	Q7Q156	Q7q156 mus musculus	214	124	7.3	311	2	Q6DN73	Q6dn73 homo sapien
142	130	7.7	253	2	Q9D8H2	Q9d8h2 m mus muscu	215	124	7.3	394	2	Q6DC16	Q6dc16 xenopus lae
143	130	7.7	285	2	Q8VE93	Q8ve93 mus musculus	216	124	7.3	597	1	SILL_PANTR	Q95l10 pan troglod
144	130	7.7	515	1	FVR1_PIG	Q9gl76 sus scrofa	217	124	7.3	623	2	Q8BY18	Q8by18 mus musculus
145	130	7.7	582	2	Q8R4B5	Q8r4b5 mus musculus	218	124	7.3	688	2	Q8QZE3	Q8qze3 mus musculus
146	130	7.7	915	2	Q8R4B3	Q8r4b3 mus musculus	219	124	7.3	697	2	Q7PMJ7	Q7pmj7 anopheles g
147	130	7.7	955	1	MDG1_HUMAN	Q8nf04 homo sapien	220	124	7.3	853	1	NCA1_BOVIN	P31836 bos taurus
148	129.5	7.7	300	2	Q68SP0	Q68sp0 mus musculus	221	124	7.3	1240	1	NFAS_HUMAN	Q94856 homo sapien
149	129.5	7.7	1040	1	AXO1_RAT	Q68ap0 mus musculus	222	124	7.3	1240	1	NFAS_MOUSE	Q810u3 mus musculus
150	129.5	7.7	1479	2	Q7KQY5	P22063 rattus norv	223	124	7.3	1240	1	NFAS_MOUSE	P97685 rattus norv
151	129.5	7.7	1482	2	Q9V4Y0	Q9v4y0 drosophila	224	124	7.3	1251	2	NFAS_RAT	Q6zq54 mus musculus
152	129.5	7.7	3375	1	UN52_CAEEL	Q85661 caenorhabdi	225	124	7.3	1366	1	ROB3_MOUSE	Q92214 mus musculus
153	129	7.6	208	2	Q80WN3	Q80wn3 mus musculus	226	124	7.3	1730	2	Q7YR07	Q7yr07 sus scrofa
154	129	7.6	324	2	Q7TMH2	Q7tmh2 mus musculus	227	123.5	7.3	316	2	Q7TPB4	Q7tpb4 rattus norv
155	129	7.6	343	2	Q8R4Y0	Q8r4y0 mus musculus	228	123.5	7.3	338	1	LAMP_CHICK	Q98919 gallus gall
156	129	7.6	433	2	Q9V644	Q9v644 drosophila	229	123.5	7.3	350	2	Q02869	Q02869 gallus gall
157	129	7.6	1427	2	Q91562	Q91562 xenopus lae	230	123.5	7.3	383	2	Q75ML9	Q75ml9 homo sapien
158	128.5	7.6	1040	1	AXO1_MOUSE	Q61330 mus musculus	231	123.5	7.3	412	2	Q63611	Q63611 rattus norv
159	128.5	7.6	2053	2	Q81ZV4	Q81zv4 homo sapien	232	123.5	7.3	454	2	Q91W54	Q91w54 mus musculus
160	128	7.6	255	2	Q9VQ64	Q9vq64 drosophila	233	123.5	7.3	521	1	CEA1_MOUSE	P31809 mus musculus
161	128	7.6	257	2	Q8R202	Q8r202 mus musculus	234	123.5	7.3	521	2	Q925P3	Q925p3 mus musculus
162	128	7.6	261	2	Q9D7L8	Q9d7l8 m mus muscu	235	123.5	7.3	605	2	Q921P2	Q921p2 mus musculus
163	128	7.6	308	2	Q68EV1	Q68ev1 xenopus lae	236	123.5	7.3	815	2	Q805B9	Q805b9 brachydanio
164	128	7.6	343	2	Q8BYS4	Q8bys4 mus musculus	237	123.5	7.3	838	2	Q8BQ96	Q8bq96 mus musculus
165	128	7.6	344	2	Q9VY33	Q9vy33 drosophila	238	123.5	7.3	838	2	Q8C4B2	Q8c4b2 mus musculus
166	128	7.6	386	2	Q8BIN0	Q8bin0 mus musculus	239	123.5	7.3	1091	1	NCA1_CHICK	P13590 gallus gall
167	128	7.6	392	2	Q7PSN2	Q7psn2 anopheles g	240	123.5	7.3	1151	2	Q9QVW5	Q9qvn5 rattus sp.
168	128	7.6	977	2	Q96RD9	Q96rd9 homo sapien	241	123.5	7.3	1256	1	NRCA_MOUSE	Q810u4 mus musculus
169	127.5	7.6	299	2	Q7Q8F3	Q7q8f3 anopheles g	242	123	7.3	265	2	Q9NGZ0	Q9ngz0 spodoptera
170	127.5	7.6	1028	2	Q6INB5	Q6inb5 xenopus lae	243	123	7.3	344	1	NTRI_HUMAN	Q9p121 homo sapien
171	127.5	7.6	1340	2	Q8NDA2	Q8nda2 homo sapien	244	123	7.3	439	2	O57349	O57349 gallus gall
172	127.5	7.6	1906	1	KMLS_CHICK	P11799 gallus gall	245	123	7.3	464	2	O6GL25	O6gl25 xenopus tro
173	127	7.5	285	2	Q8BTFO	Q8btk0 mus musculus	246	123	7.3	538	2	Q29123	Q29123 sus scrofa
174	127	7.5	337	1	OPCM_CHICK	Q98B92 gallus gall	247	123	7.3	702	2	Q89ZV8	Q89zy8 mus musculus
175	127	7.5	344	2	Q9DF61	Q9df61 gallus gall	248	123	7.3	1247	2	Q7Q0S6	Q7q0s6 anopheles g
176	127	7.5	632	2	Q6ZRK5	Q6zrk5 homo sapien	249	122.5	7.3	400	2	Q8HY16	Q8hy16 cebus apell
177	127	7.5	1249	2	Q90Z04	Q90z04 xenopus lae	250	122.5	7.3	761	1	NCA2_HUMAN	P13592 homo sapien

251	122.5	7.3	848	1	NCA1_HUMAN	P13591	homo sapien	324	119	7.0	837	2	O97137	O97137	manduca sex
252	122.5	7.3	1032	2	Q8UVD6	Q8UVD6	brachydanio	325	119	7.0	837	2	Q6RKB3	Q6RKB3	rattus norv
253	122.5	7.3	1065	1	LIG2_HUMAN	Q94898	homo sapien	326	119	7.0	1091	1	LIG1_MOUSE	LIG1_MOUSE	mus musculus
254	122.5	7.3	1284	1	NRCA_CHICK	P35331	gallus gall	327	119	7.0	1994	2	Q6ZPP2	Q6ZPP2	mus musculus
255	122.5	7.3	1419	2	Q98SW3	Q98SW3	brachydanio	328	119	7.0	2176	2	Q6V4S5	Q6V4S5	mus musculus
256	122.5	7.3	18412	2	Q7Z261	Q7Z261	brachydanio	329	119	7.0	4463	2	Q8MLD8	Q8MLD8	drosohila
257	122	7.2	240	2	Q6MG96	Q6MG96	rattus norv	330	119	7.0	9270	2	Q8MLD9	Q8MLD9	drosohila
258	122	7.2	410	2	Q6R3M2	Q6R3M2	bombyx mori	331	119	7.0	26926	2	Q10466	Q10466	homo sapien
259	122	7.2	433	2	Q6DJ83	Q6DJ83	xenopus tro	332	119	7.0	26926	2	Q8WZB3	Q8WZB3	homo sapien
260	122	7.2	1166	2	Q9QVN4	Q9QVN4	rattus sp.	333	118.5	7.0	152	2	Q8BSQ8	Q8BSQ8	mus musculus
261	121.5	7.2	320	2	Q700P8	Q700P8	anopheles g	334	118.5	7.0	262	2	Q6UXZ0	Q6UXZ0	homo sapien
262	121.5	7.2	344	1	CEA6_HUMAN	P40199	homo sapien	335	118.5	7.0	282	2	Q8VIM1	Q8VIM1	mus musculus
263	121.5	7.2	549	2	Q9NQS3	Q9NQS3	homo sapien	336	118.5	7.0	282	2	Q7TPU2	Q7TPU2	mus musculus
264	121.5	7.2	595	2	Q6ZRS5	Q6ZRS5	homo sapien	337	118.5	7.0	333	1	AMAL_DROME	AMAL_DROME	drosohila
265	121.5	7.2	1154	2	Q9QVN3	Q9QVN3	rattus sp.	338	118.5	7.0	341	2	Q7KXS2	Q7KXS2	drosohila
266	121.5	7.2	1194	2	Q6PW35	Q6PW35	rattus norv	339	118.5	7.0	413	2	Q7QBV1	Q7QBV1	anopheles g
267	121.5	7.2	1197	2	Q6PW38	Q6PW38	rattus norv	340	118.5	7.0	439	2	Q6GTU4	Q6GTU4	homo sapien
268	121.5	7.2	1198	2	Q6PW37	Q6PW37	rattus norv	341	118.5	7.0	458	2	Q83093	Q83093	rattus norv
269	121.5	7.2	1206	2	Q6PW36	Q6PW36	rattus norv	342	118.5	7.0	467	1	SIL7_HUMAN	SIL7_HUMAN	homo sapien
270	121.5	7.2	1209	2	Q6PW39	Q6PW39	rattus norv	343	118.5	7.0	510	2	Q96NY8	Q96NY8	homo sapien
271	121.5	7.2	1214	1	NRCA_RAT	P97686	rattus norv	344	118.5	7.0	519	1	ECTO_RAT	ECTO_RAT	rattus norv
272	121.5	7.2	1224	2	Q00533	Q00533	homo sapien	345	118.5	7.0	947	1	MUSK_CHICK	MUSK_CHICK	gallus gall
273	121.5	7.2	1299	2	Q6PW34	Q6PW34	rattus norv	346	118.5	7.0	1011	2	Q24273	Q24273	drosohila
274	121.5	7.2	1842	2	Q81ZY3	Q81ZY3	homo sapien	347	118.5	7.0	1476	2	Q7QJ29	Q7QJ29	anopheles g
275	121	7.2	366	2	Q6NVZ3	Q6NVZ3	homo sapien	348	118.5	7.0	1561	2	Q924D2	Q924D2	mus musculus
276	121	7.2	410	2	Q71V79	Q71V79	bombyx mori	349	118.5	7.0	1694	1	SN_MOUSE	SN_MOUSE	mus musculus
277	121	7.2	467	2	Q91VT9	Q91VT9	mus musculus	350	118	7.0	344	1	NTRI_MOUSE	NTRI_MOUSE	mus musculus
278	121	7.2	528	2	P91670	P91670	drosohila	351	118	7.0	344	2	Q8BG33	Q8BG33	m mus muscu
279	121	7.2	532	2	Q6NNU3	Q6NNU3	drosohila	352	118	7.0	359	2	Q7ZX17	Q7ZX17	anopheles lae
280	121	7.2	532	2	Q9VLF0	Q9VLF0	drosohila	353	118	7.0	391	2	Q7QJG1	Q7QJG1	anopheles g
281	121	7.2	858	2	O18466	O18466	hirudo medi	354	118	7.0	413	2	Q9VAR6	Q9VAR6	drosohila
282	121	7.2	1031	2	Q9OYM2	Q9OYM2	brachydanio	355	118	7.0	450	2	Q9VR25	Q9VR25	drosohila
283	121	7.2	1086	2	Q7QHO2	Q7QHO2	anopheles g	356	118	7.0	501	2	Q6Q147	Q6Q147	bos taurus
284	121	7.2	1155	2	Q7Q3K8	Q7Q3K8	anopheles g	357	118	7.0	540	2	Q8N029	Q8N029	homo sapien
285	121	7.2	1304	1	NRCA_HUMAN	Q92823	homo sapien	358	118	7.0	547	1	ICA3_HUMAN	ICA3_HUMAN	homo sapien
286	120.5	7.1	163	2	Q8K1H8	Q8K1H8	mus musculus	359	118	7.0	547	2	Q6PDS6	Q6PDS6	homo sapien
287	120.5	7.1	202	2	Q6NTA1	Q6NTA1	homo sapien	360	118	7.0	626	2	Q6DCH3	Q6DCH3	anopheles lae
288	120.5	7.1	265	2	Q7PDU3	Q7PDU3	anopheles g	361	118	7.0	837	1	NCM2_MOUSE	NCM2_MOUSE	mus musculus
289	120.5	7.1	358	2	Q90490	Q90490	brachydanio	362	118	7.0	1338	1	VGR1_HUMAN	VGR1_HUMAN	h vascular
290	120.5	7.1	442	1	SIL6_HUMAN	O43699	homo sapien	363	118	7.0	4824	2	Q95YM1	Q95YM1	procanbarus
291	120.5	7.1	442	2	Q9BY67	Q9BY67	homo sapien	364	118	7.0	17352	2	Q95YM2	Q95YM2	procanbarus
292	120.5	7.1	529	2	Q91V87	Q91V87	mus musculus	365	117.5	7.0	309	2	Q81YV7	Q81YV7	mus musculus
293	120.5	7.1	538	2	Q28939	Q28939	sus scrofa	366	117.5	7.0	316	2	Q8VE98	Q8VE98	mus musculus
294	120.5	7.1	549	2	Q9JLB9	Q9JLB9	mus musculus	367	117.5	7.0	338	1	LAMP_HUMAN	LAMP_HUMAN	homo sapien
295	120.5	7.1	1940	2	Q6PDN3	Q6PDN3	mus musculus	368	117.5	7.0	340	2	Q9W3N2	Q9W3N2	drosohila
296	120	7.1	304	2	Q9BPN5	Q9BPN5	caenorhabdi	369	117.5	7.0	344	2	Q13774	Q13774	homo sapien
297	120	7.1	312	2	Q6UXG6	Q6UXG6	homo sapien	370	117.5	7.0	509	2	Q920C2	Q920C2	mus musculus
298	120	7.1	410	2	Q6R3L9	Q6R3L9	bombyx mand	371	117.5	7.0	775	2	Q6PF50	Q6PF50	anopheles lae
299	120	7.1	641	2	Q86SD2	Q86SD2	ciona intes	372	117.5	7.0	800	2	Q918X3	Q918X3	brachydanio
300	120	7.1	1033	2	Q24327	Q24327	drosohila	373	117.5	7.0	858	2	Q86X47	Q86X47	homo sapien
301	120	7.1	1033	2	Q9V643	Q9V643	drosohila	374	117.5	7.0	1028	2	P97528	P97528	rattus norv
302	120	7.1	1036	1	AXO1_CHICK	P28685	gallus gall	375	117.5	7.0	1056	2	Q7ZW34	Q7ZW34	brachydanio
303	120	7.1	1093	1	LIG1_HUMAN	P28685	gallus gall	376	117.5	7.0	1092	1	NCA2_XENLA	NCA2_XENLA	anopheles lae
304	119.5	7.1	298	1	JAM2_HUMAN	P57087	homo sapien	377	117.5	7.0	1302	1	NRG_DROME	NRG_DROME	drosohila
305	119.5	7.1	298	2	Q6YNC1	Q6YNC1	homo sapien	378	117.5	7.0	1389	2	Q90Z69	Q90Z69	brachydanio
306	119.5	7.1	373	2	Q7KYP5	Q7KYP5	homo sapien	379	117.5	7.0	2008	2	Q9VEJ5	Q9VEJ5	drosohila
307	119.5	7.1	463	1	SIL9_HUMAN	Q9V336	homo sapien	380	117.5	7.0	2046	2	Q7KSE9	Q7KSE9	drosohila
308	119.5	7.1	464	1	O16170	O16170	homo sapien	381	117.5	7.0	8625	2	Q86GD6	Q86GD6	procanbarus
309	119.5	7.1	468	2	Q96CA7	Q96CA7	homo sapien	382	117	6.9	351	2	Q8WEB2	Q8WEB2	branchiosteo
310	119.5	7.1	526	1	CEA1_HUMAN	P13688	homo sapien	383	117	6.9	401	2	Q7PSS8	Q7PSS8	anopheles g
311	119.5	7.1	529	2	Q7TQM3	Q7TQM3	rattus norv	384	117	6.9	448	2	Q9JHL7	Q9JHL7	rattus norv
312	119.5	7.1	549	2	Q9D006	Q9D006	mus musculus	385	117	6.9	459	2	Q9JHL6	Q9JHL6	rattus norv
313	119.5	7.1	725	1	NCA2_MOUSE	P13594	mus musculus	386	117	6.9	595	1	SILL_HUMAN	SILL_HUMAN	homo sapien
314	119.5	7.1	725	2	Q73634	Q73634	anopheles lae	387	117	6.9	739	1	VCA1_RAT	VCA1_RAT	rattus norv
315	119.5	7.1	1115	1	NCA1_MOUSE	P13595	mus musculus	388	117	6.9	858	1	NCA1_RAT	NCA1_RAT	rattus norv
316	119.5	7.1	1643	2	Q7QGT8	Q7QGT8	anopheles g	389	117	6.9	1062	2	Q8BK93	Q8BK93	mus musculus
317	119.5	7.1	1950	2	Q8QYN8	Q8QYN8	mus musculus	390	117	6.9	1445	2	Q63155	Q63155	rattus norv
318	119	7.0	334	2	O02870	O02870	gallus gall	391	117	6.9	1447	1	DCC_HUMAN	DCC_HUMAN	homo sapien
319	119	7.0	337	2	P97268	P97268	cavia porce	392	117	6.9	2164	2	Q9IAR9	Q9IAR9	gallus gall
320	119	7.0	410	2	Q6R3M0	Q6R3M0	bombyx mori	393	116.5	6.9	285	2	Q7PNI4	Q7PNI4	anopheles g
321	119	7.0	467	2	Q8C6F2	Q8C6F2	mus musculus	394	116.5	6.9	338	1	LAMP_RAT	LAMP_RAT	rattus norv
322	119	7.0	727	2	Q6RKB2	Q6RKB2	rattus norv	395	116.5	6.9	341	1	LAMP_MOUSE	LAMP_MOUSE	mus musculus
323	119	7.0	779	2	O97136	O97136	manduca sex	396	116.5	6.9	378	2	Q66MN4	Q66MN4	petromyzon

397	116.5	6.9	445	2	Q8K3T6	Q8K3t6 mus musculus
398	116.5	6.9	554	2	Q8W4R3	Q8W4r3 drosophila
399	116.5	6.9	806	1	CEK2_CHICK	P18460 gallus gall
400	116.5	6.9	899	2	Q7PQM9	Q7Pqm9 anopheles g
401	116.5	6.9	972	2	Q26614	Q26614 strongyloce
402	116.5	6.9	1040	1	AX01_HUMAN	Q02246 homo sapien
403	116.5	6.9	1051	1	PTK7_CHICK	Q91048 gallus gall
404	116.5	6.9	1598	2	Q8P2I4	Q8P2i4 homo sapien
405	116	6.9	237	2	Q6DQX5	Q6dqx5 oryctolagus
406	116	6.9	344	1	NTRI_RAT	Q62718 rattus norv
407	116	6.9	344	1	Q6B0I4	Q6B0i4 homo sapien
408	116	6.9	1027	2	Q9OW79	Q9ow79 gallus gall
409	116	6.9	1103	2	Q6ONE3	Q6onf3 canis fami
410	116	6.9	1447	1	DCC_MOUSE	P70211 mus musculu
411	116	6.9	1709	1	SN_HUMAN	Q9Bz22 homo sapien
412	116	6.9	6658	2	Q76281	Q76281 drosophila
413	116	6.9	8647	2	Q7KQP5	Q7kqp5 drosophila
414	116	6.9	8648	2	Q7KQP6	Q7kqp6 drosophila
415	116	6.9	8930	2	Q7KQP7	Q7kqp7 drosophila
416	116	6.9	8943	2	Q9V4F7	Q9v4f7 drosophila
417	115.5	6.8	295	2	Q9QYL6	Q9qyl6 mus musculu
418	115.5	6.8	295	2	Q9Z2H8	Q9z2h8 mus musculu
419	115.5	6.8	309	1	CD86_MOUSE	P42082 mus musculu
420	115.5	6.8	314	2	Q61238	Q61238 homo sapien
421	115.5	6.8	356	2	Q64381	Q64381 mus musculu
422	115.5	6.8	390	2	Q8P500	Q8P500 rattus norv
423	115.5	6.8	445	2	Q8R4L1	Q8r4l1 mus musculu
424	115.5	6.8	520	2	Q9Z5P2	Q9z5p2 mus musculu
425	115.5	6.8	725	2	Q73633	Q73633 xenopus lae
426	115.5	6.8	738	1	PEC1_HUMAN	P16284 homo sapien
427	115.5	6.8	1252	2	Q96DN3	Q96dn3 homo sapien
428	115.5	6.8	1255	2	Q723Z9	Q723z9 homo sapien
429	115.5	6.8	1255	2	Q71QL8	Q71ql8 pan troglod
430	115.5	6.8	1257	1	CAML_HUMAN	P32004 homo sapien
431	115.5	6.8	1493	1	NEO1_MOUSE	P97798 mus musculu
432	115.5	6.8	4001	2	Q9NL2P7	Q9nl2p7 drosophila
433	115.5	6.8	4796	2	Q9NL88	Q9nl88 drosophila
434	115.5	6.8	4796	2	Q9W055	Q9w055 drosophila
435	115.5	6.8	16215	2	Q9NFS3	Q9nfs3 drosophila
436	115.5	6.8	17903	2	Q7RTL4	Q7rtl4 drosophila
437	115.5	6.8	18074	2	Q917U4	Q917u4 drosophila
438	115	6.8	217	2	Q6KGN0	Q6kgn0 bacterioph
439	115	6.8	237	2	Q8CG82	Q8cg82 mus musculu
440	115	6.8	294	2	Q6KGN1	Q6kgn1 bacterioph
441	115	6.8	306	2	Q9DYL4	Q9dy14 mus musculu
442	115	6.8	316	2	Q8WPB3	Q8wpb3 drosophila
443	115	6.8	399	2	Q8N7P2	Q8n7p2 homo sapien
444	115	6.8	424	2	Q8C6W0	Q8c6w0 mus musculu
445	115	6.8	602	2	Q8GYJ9	Q8gyj9 homo sapien
446	115	6.8	650	2	Q8N8A4	Q8na84 homo sapien
447	115	6.8	731	2	Q8SP16	Q8spi6 macropus eu
448	115	6.8	837	2	Q7Z7F2	Q7z7f2 homo sapien
449	115	6.8	1125	2	Q7QEC1	Q7qec1 anopheles g
450	115	6.8	1209	2	P70232	P70232 mus musculu
451	115	6.8	1302	1	VGR2_BRARE	Q8axb3 brachydanio
452	115	6.8	1723	2	Q8CHE2	Q8chb2 mus musculu
453	115	6.8	2403	2	Q8MLD5	Q8mld5 drosophila
454	115	6.8	7210	2	Q9V7G8	Q9v7g8 drosophila
455	114.5	6.8	163	2	Q9NVJ5	Q9nvj5 homo sapien
456	114.5	6.8	345	2	Q6QM08	Q6qm08 xenopus lae
457	114.5	6.8	363	2	Q6NV41	Q6nv41 brachydanio
458	114.5	6.8	412	2	Q8HY14	Q8hy14 oryctolagus
459	114.5	6.8	458	2	Q61351	Q61351 mus musculu
460	114.5	6.8	484	2	Q6BE00	Q6be00 xenopus lae
461	114.5	6.8	487	2	Q7PGL7	Q7pgl7 anopheles g
462	114.5	6.8	499	2	Q9BXN7	Q9bxn7 homo sapien
463	114.5	6.8	499	1	SIL8_HUMAN	Q9ny24 homo sapien
464	114.5	6.8	504	2	Q8N441	Q8n441 homo sapien
465	114.5	6.8	504	2	Q9H4D7	Q9h4d7 homo sapien
466	114.5	6.8	521	2	Q61352	Q61352 mus musculu
467	114.5	6.8	538	1	PVR2_HUMAN	Q92692 homo sapien
468	114.5	6.8	551	2	Q8MSN7	Q8msn7 drosophila
469	114.5	6.8	719	2	Q661V0	Q661v0 xenopus lae

Q9W4t9	drosophila	Q9W4T9	2	956	6.8	114.5	470
Q8n9y3	drosophila	Q8N9Y3	2	959	6.8	114.5	471
Q97174	drosophila	Q97174	2	975	6.8	114.5	472
P16170	xenopus lae	NCAL_XENLA	1	1088	6.8	114.5	473
Q84155	caenorhabdi	Q84155	2	1415	6.8	114.5	474
Q8wz53	homo sapien	Q8WZ53	2	5604	6.8	114.5	475
Q75296	homo sapien	Q75296	2	235	6.8	114	476
Q6dfv2	mus musculu	Q6DFY2	2	337	6.8	114	477
Q86kx2	xenopus lae	Q86KX2	2	390	6.8	114	478
Q8hy15	lemur catta	Q8HY15	2	403	6.8	114	479
Q9j1b7	mus musculu	Q9JL87	2	438	6.8	114	480
Q9dbp8	mus musculu	Q9DBP8	2	483	6.8	114	481
Q8ced8	mus musculu	Q8CED8	2	508	6.8	114	482
Q8r007	mus musculu	Q8R007	2	508	6.8	114	483
Q9j1b8	mus musculu	Q9JL88	2	510	6.8	114	484
Q28125	bos taurus	IC3A_BOVIN	1	544	6.8	114	485
Q9vct4	drosophila	Q9VCT4	2	545	6.8	114	486
Q8ip70	drosophila	Q8IP70	2	672	6.8	114	487
Q90ym0	brachydanio	Q90YMO	2	795	6.8	114	488
Q8c6x1	mus musculu	Q8C6X1	2	1028	6.8	114	489
Q9jmb8	mus musculu	Q9JMB8	2	1028	6.8	114	490
Q8ct37	homo sapien	Q8CT37	2	1045	6.8	114	491
Q8txi8	caenorhabdi	Q8TXI8	2	1073	6.8	114	492
Q6gqb1	xenopus lae	Q6GQB1	2	1177	6.8	114	493
Q96kf5	homo sapien	Q96KF5	2	1320	6.8	114	494
Q8ctc9	homo sapien	Q8CTC9	2	1320	6.8	114	495
Q7q916	anopheles g	Q7Q916	2	1735	6.8	114	496
Q7fj18	anopheles g	Q7FJ18	2	232	6.7	113.5	497
Q9nx42	homo sapien	Q9NX42	2	284	6.7	113.5	498
Q6pj56	homo sapien	Q6PJ56	2	296	6.7	113.5	499
Q96iq7	homo sapien	Q96IQ7	2	327	6.7	113.5	500
Q7psh7	anopheles g	Q7PSH7	2	382	6.7	113.5	501
Q7z728	homo sapien	Q7Z728	2	499	6.7	113.5	502
Q96k15	homo sapien	Q96K15	2	510	6.7	113.5	503
P32507	mus musculu	PVR2_MOUSE	1	530	6.7	113.5	504
Q80xj5	mus musculu	Q80XJ5	2	530	6.7	113.5	505
Q8bz76	m mus muscu	Q8BZ76	2	754	6.7	113.5	506
Q7qey8	anopheles g	Q7QEY8	2	771	6.7	113.5	507
Q9zy7	mus musculu	Q9ZY7	2	875	6.7	113.5	508
Q967x6	drosophila	Q967X6	2	879	6.7	113.5	509
Q9vb35	drosophila	Q9VB35	2	939	6.7	113.5	510
Q9xt41	cercopithec	Q9XT41	2	1248	6.7	113.5	511
Q7yql7	pongo pygma	Q7YQL7	2	1255	6.7	113.5	512
Q96ms0	homo sapien	Q96MS0	1	1385	6.7	113.5	513
Q8uud7	xenopus lae	Q8UUD7	2	1614	6.7	113.5	514
Q98sw4	brachydanio	Q98SW4	2	1675	6.7	113.5	515
Q15762	homo sapien	C226_HUMAN	1	336	6.7	113	516
Q7q807	anopheles g	Q7Q807	2	340	6.7	113	517
Q9nw07	homo sapien	Q9NW07	2	538	6.7	113	518
Q8p5h3	mus musculu	Q8P5H3	2	1109	6.7	113	519
Q89902	fugu rubrip	CAML_FUGRU	1	1277	6.7	113	520
Q7kyn5	homo sapien	Q7KYN5	2	3100	6.7	113	521
Q15598	homo sapien	Q15598	2	4650	6.7	113	522
Q7pyg1	anopheles g	Q7PYG1	2	226	6.7	112.5	523
Q95t89	drosophila	Q95T89	2	244	6.7	112.5	524
Q7z3w6	homo sapien	Q7Z3W6	2	338	6.7	112.5	525
Q14982	homo sapien	Q14982	1	345	6.7	112.5	526
Q9jhl1	rattus norv	Q9JHQ1	2	362	6.7	112.5	527
Q46147	onchocerca	Q46147	2	363	6.7	112.5	528
Q08835	cercopithec	Q08835	2	401	6.7	112.5	529
Q6zmd4	homo sapien	Q6ZMD4	2	440	6.7	112.5	530
Q15223	homo sapien	PVR1_HUMAN	1	517	6.7	112.5	531
Q8f6b2	vibriophaga	Q8F6B2	2	743	6.7	112.5	532
Q61563	mus musculu	Q61563	2	782	6.7	112.5	533
Q95695	rattus norv	CAML_RAT	1	1259	6.7	112.5	534
Q6pgj3	mus musculu	Q6PGJ3	2	1342	6.7	112.5	535
Q9gpp6	drosophila	Q9GPP6	2	1342	6.7	112.5	536
Q9vpz7	drosophila	Q9VPZ7	2	1342	6.7	112.5	537
P16f21	drosophila	LAR_DRONE	1	2029	6.7	112.5	538
Q9vi88	drosophila	Q9VI88	2	2029	6.7	112.5	539
Q9ulm1	drosophila	Q9ULM1	2	2224	6.7	112.5	540
Q8wp94	drosophila	Q8WP94	2	316	6.6	112	541
Q7z3b1	homo sapien	NBGR_HUMAN	1	352	6.6	112	542

543	112	387	2	Q64JA4	Q64ja4 pan troglod	616	109.5	6.5	1465	2	Q7TQGS	Q7tqgs: mus musculus
544	112	456	2	Q8R5M8	Q8r5m8 mus musculus	617	109	6.5	263	2	Q7TPW5	Q7tpw5 mus musculus
545	112	476	2	Q6AYP5	Q6ayp5 rattus norv	618	109	6.5	283	2	Q7TPH5	Q7tpH5 mus musculus
546	112	590	2	Q6P4T5	Q6p4t5 mus musculus	619	109	6.5	283	2	Q7TSP5	Q7tap5 mus musculus
547	112	913	2	Q8T3E5	Q8t3e5 caenorhabdi	620	109	6.5	292	2	Q6UY47	Q6uy47 homo sapien
548	112	928	2	Q13128	Q19128 caenorhabdi	621	109	6.5	316	2	Q8WP58	Q8wp58 drosophila
549	112	946	2	Q07155	Q07153 torpedo cal	622	109	6.5	328	2	O88775	O88775 rattus norv
550	112	955	2	Q8MQ86	Q8mq86 caenorhabdi	623	109	6.5	336	2	O46551	O46551 hylobates s
551	112	1212	2	Q95TGO	Q95tgo drosophila	624	109	6.5	345	2	Q7PVP1	Q7pvp1 anopheles s
552	112	1269	2	Q01632	Q01632 caenorhabdi	625	109	6.5	437	2	Q8IZP8	Q8izp8 homo sapien
553	112	1273	2	O44928	O44928 caenorhabdi	626	109	6.5	547	1	CB19_MOUSE	P25918 mus musculus
554	112	1386	2	O8TI72	O8ti72 methanosarc	627	109	6.5	562	2	Q8BSM2	Q6ynr7 brachydanio
555	111.5	267	2	Q8NC05	Q8nc05 homo sapien	628	109	6.5	640	2	Q8BSM2	Q8bem2 mus musculus
556	111.5	390	2	Q95K13	Q95k13 macaca fasc	629	109	6.5	646	1	MU18_HUMAN	P43121 homo sapien
557	111.5	400	1	HPL4_MOUSE	Q80wm4 mus musculus	630	109	6.5	646	2	Q6PHR3	Q6phr3 homo sapien
558	111.5	508	2	Q96LA5	Q96la5 homo sapien	631	109	6.5	1106	2	O8WX93	O8wx93 homo sapien
559	111.5	1060	2	Q90Z13	Q9qz13 rattus norv	632	108.5	6.4	345	1	OPCM_BOVIN	P11834 bos taurus
560	111	252	2	Q8WWT6	Q8wwt6 homo sapien	633	108.5	6.4	348	1	NEGR_MOUSE	Q80z24 mus musculus
561	111	283	2	Q8K091	Q8k091 mus musculus	634	108.5	6.4	348	1	NEGR_RAT	Q9z0j8 rattus norv
562	111	421	2	Q7PV30	Q7pv30 anopheles g	635	108.5	6.4	416	2	Q7M048	Q7m048 rattus norv
563	111	421	2	Q7QLK4	Q7qlk4 anopheles g	636	108.5	6.4	495	2	Q9HCY1	Q9hcy1 homo sapien
564	111	474	2	Q7PKE3	Q7pke3 anopheles g	637	108.5	6.4	705	2	Q8CBD3	Q8cbd3 mus musculus
565	111	527	2	Q9ERF7	Q9erf7 cricetus	638	108.5	6.4	878	2	Q9GV22	Q9gv22 mytilus gal
566	111	769	2	Q8NI15	Q8ni15 homo sapien	639	108.5	6.4	898	2	Q9VZ26	Q9vz26 mus musculus
567	111	806	1	FGK3_HUMAN	P22607 homo sapien	640	108.5	6.4	1028	2	Q9UQ52	Q9uq52 homo sapien
568	111	807	2	Q6NY23	Q6ny23 brachydanio	641	108.5	6.4	1114	2	Q9BWW1	Q9bww1 homo sapien
569	111	960	2	Q7PV74	Q7pv74 anopheles g	642	108.5	6.4	1115	2	Q6UXJ5	Q6uxj5 homo sapien
570	111	1026	2	Q8IWW2	Q8iww2 homo sapien	643	108.5	6.4	1328	2	O21043	O21043 caenorhabdi
571	111	1228	2	Q8WRA3	Q8wra3 drosophila	644	108.5	6.4	2000	2	O87791	O87791 oryctolagus
572	111	1325	2	Q8GBD5	Q8gbd5 drosophila	645	108.5	6.4	2169	2	Q8AV58	Q8av58 gallus gall
573	111	1335	2	Q9V787	Q9v787 drosophila	646	108.5	6.4	4203	2	Q965G2	Q965g2 caenorhabdi
574	111	6632	1	UN89_CABEL	O01761 caenorhabdi	647	108.5	6.4	4219	2	Q9NL87	Q9nl87 caenorhabdi
575	110.5	8081	2	Q72120	Q72120 caenorhabdi	648	108.5	6.4	4369	2	Q8MXD7	Q8mxd7 caenorhabdi
576	110.5	370	2	Q6WZQ3	Q6wzq3 homo sapien	649	108.5	6.4	4447	2	Q8MXD8	Q8mxd8 caenorhabdi
577	110.5	390	2	Q95NP7	Q95np7 homo sapien	650	108.5	6.4	4488	2	Q9TXK2	Q9txk2 caenorhabdi
578	110.5	422	2	Q96FP3	Q96fp3 homo sapien	651	108.5	6.4	4736	2	Q7YT99	Q7yt99 mytilus gal
579	110.5	513	2	O00481	Q96pj3 homo sapien	652	108	6.4	173	2	Q9JKD5	Q9jkd5 rattus norv
580	110.5	515	2	Q96PJ5	Q96pj5 homo sapien	653	108	6.4	286	2	O46535	O46535 bos taurus
581	110.5	515	2	Q96PJ5	Q96pj5 homo sapien	654	108	6.4	325	2	O95791	O95791 homo sapien
582	110.5	515	2	Q96RE0	Q96re0 homo sapien	655	108	6.4	325	2	Q8HW98	Q8hw98 mus musculus
583	110.5	544	2	Q6UXI8	Q6uxi8 homo sapien	656	108	6.4	329	2	Q8N225	Q8n225 homo sapien
584	110.5	816	2	Q91285	Q91285 pleurodeles	657	108	6.4	332	2	Q819N2	Q819n2 brachiosco
585	110.5	1070	1	PTK7_HUMAN	Q13308 homo sapien	658	108	6.4	332	2	Q90Z71	Q90z71 brachydanio
586	110.5	1189	2	Q9P2J2	Q9p2j2 homo sapien	659	108	6.4	509	2	Q9EQY5	Q9eqy5 m mman-g pr
587	110.5	1889	2	Q7Q0X2	Q7q0x2 anopheles g	660	108	6.4	526	1	BUTY_BOVIN	P18892 bos taurus
588	110	218	2	Q6ZMC6	Q6zmc6 homo sapien	661	108	6.4	614	2	Q6DDQ7	Q6ddq7 xenopus lae
589	110	316	2	Q9EXR1	Q9bxr1 homo sapien	662	108	6.4	868	1	MUSK_RAT	Q62838 rattus norv
590	110	483	2	Q78X76	Q78x76 brachydanio	663	108	6.4	1099	2	P97527	P97527 rattus norv
591	110	636	2	Q22040	Q22040 caenorhabdi	664	108	6.4	1150	2	Q8BS24	Q8be24 mus musculus
592	110	741	2	Q8RCF4	Q8rcf4 vibrionphage	665	108	6.4	1244	2	Q69YJ3	Q69yj3 homo sapien
593	110	815	2	Q8AYP3	Q8ayp3 brachydanio	666	108	6.4	2325	2	Q9N3X8	Q9n3x8 caenorhabdi
594	110	837	1	Q8NFA5	Q8nfa5 homo sapien	667	107.5	6.4	323	2	Q9BDB8	Q9bdb8 cercocebus
595	110	837	1	NCW2_HUMAN	O15394 homo sapien	668	107.5	6.4	350	2	Q99420	Q99420 homo sapien
596	110	917	1	ICA5_MOUSE	Q60e25 mus musculus	669	107.5	6.4	350	2	Q819N1	Q819n1 brachiosco
597	110	1070	2	Q61Q54	Q61q54 homo sapien	670	107.5	6.4	366	2	Q8N759	Q8n759 homo sapien
598	110	1097	1	PGDR_RAT	Q05030 rattus norv	671	107.5	6.4	402	1	HPL4_HUMAN	Q86uw8 homo sapien
599	110	1378	1	ROB2_HUMAN	Q9hck4 homo sapien	672	107.5	6.4	428	2	Q96PJ6	Q96pj6 homo sapien
600	110	1391	2	O8N3L4	Q8n3l4 homo sapien	673	107.5	6.4	429	2	Q96LA6	Q96la6 homo sapien
601	110	1880	2	O18465	O18465 hirudo medi	674	107.5	6.4	531	2	Q659F2	Q659f2 homo sapien
602	110	2389	2	Q6BEQ6	Q6beq6 caenorhabdi	675	107.5	6.4	605	2	Q6GNL9	Q6gnl9 xenopus lae
603	109.5	323	2	Q9BDM2	Q9bdm2 cercopithe	676	107.5	6.4	659	2	Q6ZNM1	Q6znm1 homo sapien
604	109.5	333	1	C226_MOUSE	Q8k4f0 mus musculus	677	107.5	6.4	660	2	Q7Z681	Q7z681 homo sapien
605	109.5	334	2	O819N0	Q819n0 brachiosco	678	107.5	6.4	1026	2	Q62845	Q62845 rattus norv
606	109.5	345	1	OPCM_RAT	P32736 rattus norv	679	107.5	6.4	1225	2	O6GP61	O6gp61 xenopus lae
607	109.5	349	1	CEA8_HUMAN	P31997 homo sapien	680	107.5	6.4	1235	2	Q95428	Q95428 homo sapien
608	109.5	383	2	Q7QBC5	Q7qbc5 anopheles g	681	107	6.3	199	2	Q8ND10	Q8nd10 homo sapien
609	109.5	430	2	Q7QGS8	Q7qgs8 anopheles g	682	107	6.3	332	2	Q6UXG3	Q6uxg3 homo sapien
610	109.5	435	2	Q8N3J6	Q8n3j6 homo sapien	683	107	6.3	437	2	Q8NFS6	Q8nfs6 homo sapien
611	109.5	477	2	Q6UXJ4	Q6uxj4 homo sapien	684	107	6.3	442	2	O6KAT6	O6kat6 mus musculus
612	109.5	526	1	BUTY_HUMAN	O13410 homo sapien	685	107	6.3	449	2	Q8UE16	Q8ue16 homo sapien
613	109.5	526	2	Q9H458	Q9h458 homo sapien	686	107	6.3	515	1	PVR1_MOUSE	Q9jkl6 mus musculus
614	109.5	595	2	Q90720	Q90720 gallus gall	687	107	6.3	515	2	Q6P9M9	Q6p9m9 mus musculus
615	109.5	1336	1	VGR1_RAT	P53767 rattus norv	688	107	6.3	606	2	Q9ESS8	Q9ess8 rattus norv

689	107	6.3	622	2	Q9JKB2	Q91kb2 mus musculus
690	107	6.3	648	2	Q9EPF2	Q9epf2 rattus norv
691	107	6.3	773	1	PIGR_RABIT	P01832 oryctolagus
692	107	6.3	817	2	Q8UG38	Q81j38 brachydanio
693	107	6.3	2558	2	Q6NR91	Q6nr91 drosophila
694	107	6.3	19066	2	Q80R18	Q80r18 brachydanio
695	106.5	6.3	289	2	Q9QYL5	Q9qyl5 mus musculus
696	106.5	6.3	304	2	Q9CVA4	Q9cva4 mus musculus
697	106.5	6.3	354	1	HP1L_HORSE	Q28381 equus caball
698	106.5	6.3	437	2	Q86FV1	Q86fy1 homo sapien
699	106.5	6.3	546	2	Q80X70	Q80x70 mus musculus
700	106.5	6.3	548	2	Q99NB3	Q99nb3 mus musculus
701	106.5	6.3	719	2	Q9U4G1	Q9u4g1 drosophila
702	106.5	6.3	793	2	Q70246	Q70246 mus musculus
703	106.5	6.3	813	2	Q8BQC3	Q8bqc3 mus musculus
704	106.5	6.3	1227	2	Q21038	Q21038 caenorhabdi
705	106.5	6.3	1232	2	Q90284	Q90284 carassius a
706	106	6.3	1306	2	Q6UXI2	Q6uxi2 homo sapien
707	106	6.3	317	2	Q8BEK4	Q8bek4 cowpox viru
708	106	6.3	381	2	Q8P4B1	Q8p4b1 mus musculus
709	106	6.3	407	2	Q9D2J4	Q9d2j4 mus musculus
710	106	6.3	582	2	Q80WN2	Q80wn2 mus musculus
711	106	6.3	595	2	Q68SN8	Q68sn8 mus musculus
712	106	6.3	907	1	Q9NEG0	Q9neg0 drosophila
713	106	6.3	919	1	UNC5_CAEEL	Q26261 caenorhabdi
714	106	6.3	1040	1	EG15_CAEEL	Q10656 caenorhabdi
715	106	6.3	1051	2	Q7JL68	Q7jl68 caenorhabdi
716	106	6.3	1081	2	Q69ZT7	Q69zt7 mus musculus
717	106	6.3	1096	2	Q8MQ14	Q8mq14 caenorhabdi
718	106	6.3	1098	1	PGDR_MOUSE	P05622 mus musculus
719	106	6.3	1109	2	Q8CE91	Q8ce91 mus musculus
720	106	6.3	1110	2	Q8CE73	Q8ce73 mus musculus
721	106	6.3	1409	2	Q8J127	Q8j127 brachydanio
722	106	6.3	1409	2	Q80IM2	Q80im2 brachydanio
723	106	6.3	1428	2	Q8AY67	Q8ay67 brachydanio
724	106	6.3	1474	2	Q8T4M0	Q8t4m0 drosophila
725	106	6.3	1509	2	Q81PG1	Q81pg1 drosophila
726	106	6.3	1509	2	Q95P10	Q95p10 drosophila
727	106	6.3	1914	1	KMLS_HUMAN	Q15746 homo sapien
728	106	6.3	1914	2	Q7Z4J0	Q7z4j0 homo sapien
729	106	6.3	2222	2	Q70EG7	Q70eg7 anopheles g
730	105.5	6.2	318	2	Q8AYZ8	Q8ayz8 variola vir
731	105.5	6.2	318	2	Q8BEI5	Q8bei5 variola vir
732	105.5	6.2	328	2	Q92109	Q92109 mus musculus
733	105.5	6.2	394	2	Q6UXG0	Q6uxg0 homo sapien
734	105.5	6.2	408	2	Q8K094	Q8k094 m hypocheti
735	105.5	6.2	408	2	Q91WP1	Q91wp1 mus musculus
736	105.5	6.2	408	2	Q8BVF6	Q8bvf6 mus musculus
737	105.5	6.2	485	1	PODX_RAT	Q9wtq2 rattus norv
738	105.5	6.2	525	2	Q7P2S8	Q7pz88 anopheles g
739	105.5	6.2	527	2	Q6ZTR2	Q6ztr2 homo sapien
740	105.5	6.2	626	1	MAG_RAT	P07722 rattus norv
741	105.5	6.2	729	2	Q63827	Q63827 rattus norv
742	105.5	6.2	731	2	Q8CFK8	Q8cfk8 mus musculus
743	105.5	6.2	733	2	Q60830	Q60830 mus musculus
744	105.5	6.2	733	2	Q80T10	Q80t10 mus musculus
745	105.5	6.2	789	2	Q7PME2	Q7pme2 anopheles g
746	105.5	6.2	820	2	Q8CIM9	Q8cim9 mus musculus
747	105.5	6.2	822	1	FGRI_HUMAN	P11362 homo sapien
748	105.5	6.2	822	1	FGRI_MOUSE	P16092 mus musculus
749	105.5	6.2	822	1	FGRI_RAT	Q04589 rattus norv
750	105.5	6.2	822	2	Q60818	Q60818 mus musculus
751	105.5	6.2	869	1	MUSK_HUMAN	Q15146 homo sapien
752	105.5	6.2	987	2	Q71ZM6	Q71z88 caenorhabdi
753	105.5	6.2	1036	2	Q88WJ3	Q88wj3 drosophila
754	105.5	6.2	1102	2	Q923W7	Q923w7 mus musculus
755	105.5	6.2	1390	2	Q9VN14	Q9vn14 drosophila
756	105	6.2	145	2	Q9MZE4	Q9mze4 macaca mula
757	105	6.2	232	2	Q8BZ74	Q8bz74 mus musculus
758	105	6.2	208	2	Q9VT83	Q9vt83 drosophila
759	105	6.2	509	2	Q91YK7	Q91yk7 mus musculus
760	105	6.2	533	2	Q8NCB6	Q8ncb6 homo sapien
761	105	6.2	622	2	Q9ESS5	Q9ess5 mus musculus

Q9r069	mus musculus	Q9r069	2	622	105	6.2	762	105	6.2	Q9R069	Q9r069 mus musculus
Q95812	homo sapien	Q95812	2	646	105	6.2	763	105	6.2	Q95812	Q95812 homo sapien
Q99k86	mus musculus	Q99k86	2	650	105	6.2	764	105	6.2	Q99K86	Q99k86 mus musculus
Q8mqz9	drosophila	Q8mqz9	2	662	105	6.2	765	105	6.2	Q8MQZ9	Q8mqz9 drosophila
Q8vgd0	drosophila	Q8vgd0	2	662	105	6.2	766	105	6.2	Q8VGD0	Q8vgd0 drosophila
Q865f2	oryctolagus	Q865f2	2	739	105	6.2	767	105	6.2	Q865F2	Q865f2 oryctolagus
Q9jhx9	rattus norv	Q9jhx9	2	800	105	6.2	768	105	6.2	Q9JHX9	Q9jhx9 rattus norv
Q9pe96	xenopus lae	Q9pe96	2	810	105	6.2	769	105	6.2	Q9PE96	Q9pe96 xenopus lae
Q8gk3	xenopus lae	Q8gk3	2	828	105	6.2	770	105	6.2	Q8GK3	Q8gk3 xenopus lae
Q977y2	thermoplasma	Q977y2	2	998	105	6.2	771	105	6.2	Q977Y2	Q977y2 thermoplasma
Q8mer5	drosophila	Q8mer5	2	1066	105	6.2	772	105	6.2	Q8MSR5	Q8mer5 drosophila
Q7kti7	drosophila	Q7kti7	2	1215	105	6.2	773	105	6.2	Q7KTI7	Q7kti7 drosophila
Q8t9f6	drosophila	Q8t9f6	2	1461	105	6.2	774	105	6.2	Q8TFJ6	Q8t9f6 drosophila
Q7kti8	drosophila	Q7kti8	2	1503	105	6.2	775	105	6.2	Q7KTI8	Q7kti8 drosophila
Q9vlq8	drosophila	Q9vlq8	2	1509	105	6.2	776	105	6.2	Q9VLQ8	Q9vlq8 drosophila
Q9irv7	drosophila	Q9irv7	2	3215	105	6.2	777	105	6.2	Q9IRV7	Q9irv7 drosophila
Q9irv9	drosophila	Q9irv9	2	4117	105	6.2	778	105	6.2	Q9IRV9	Q9irv9 drosophila
Q9w4y4	drosophila	Q9w4y4	2	4179	105	6.2	779	105	6.2	Q9W4Y4	Q9w4y4 drosophila
Q8mpn3	drosophila	Q8mpn3	2	4223	105	6.2	780	105	6.2	Q8MPN3	Q8mpn3 drosophila
Q8irv8	drosophila	Q8irv8	2	4228	105	6.2	781	105	6.2	Q8IRV8	Q8irv8 drosophila
Q8t103	bombyx mori	Q8t103	2	4816	105	6.2	782	105	6.2	Q8T103	Q8t103 bombyx mori
Q02280	caenorhabdi	Q02280	2	265	104.5	6.2	783	104.5	6.2	Q02280	Q02280 caenorhabdi
Q8mre6	drosophila	Q8mre6	2	360	104.5	6.2	784	104.5	6.2	Q8MRE6	Q8mre6 drosophila
Q8bxj7	m mus muscu	Q8bxj7	2	395	104.5	6.2	785	104.5	6.2	Q8BXJ7	Q8bxj7 m mus muscu
Q8bzp4	mus musculus	Q8bzp4	2	395	104.5	6.2	786	104.5	6.2	Q8BZP4	Q8bzp4 mus musculus
P32506	cercopithec	P32506	2	417	104.5	6.2	787	104.5	6.2	P32506	P32506 cercopithec
Q8n4f1	homo sapien	Q8n4f1	2	430	104.5	6.2	788	104.5	6.2	Q8NF1	Q8n4f1 homo sapien
Q8k2h7	mus musculus	Q8k2h7	2	523	104.5	6.2	789	104.5	6.2	Q8K2H7	Q8k2h7 mus musculus
Q9gyq7	mus musculus	Q9gyq7	2	538	104.5	6.2	790	104.5	6.2	Q9GYQ7	Q9gyq7 mus musculus
Q95n25	bos taurus	Q95n25	2	582	104.5	6.2	791	104.5	6.2	Q95N25	Q95n25 bos taurus
Q6gnb3	xenopus lae	Q6gnb3	2	722	104.5	6.2	792	104.5	6.2	Q6GNB3	Q6gnb3 xenopus lae
Q9p232	homo sapien	Q9p232	2	920	104.5	6.2	793	104.5	6.2	Q9P232	Q9p232 homo sapien
Q8kam5	mus musculus	Q8kam5	2	931	104.5	6.2	794	104.5	6.2	Q8KAM5	Q8kam5 mus musculus
Q8azb0	mus musculus	Q8azb0	2	1109	104.5	6.2	795	104.5	6.2	Q8AZB0	Q8azb0 mus musculus
Q75339	homo sapien	Q75339	2	1184	104.5	6.2	796	104.5	6.2	Q75339	Q75339 homo sapien
Q6uw99	homo sapien	Q6uw99	2	1184	104.5	6.2	797	104.5	6.2	Q6UW99	Q6uw99 homo sapien
Q7prk4	anopheles g	Q7prk4	2	1185	104.5	6.2	798	104.5	6.2	Q7PRK4	Q7prk4 anopheles g
P97603	rattus norv	P97603	2	1377	104.5	6.2	799	104.5	6.2	P97603	P97603 rattus norv
Q92859	homo sapien	Q92859	2	1461	104.5	6.2	800	104.5	6.2	Q92859	Q92859 homo sapien
Q87394	drosophila	Q87394	2	2222	104.5	6.2	801	104.5	6.2	Q87394	Q87394 drosophila
Q86bq7	drosophila	Q86bq7	2	2230	104.5	6.2	802	104.5	6.2	Q86BQ7	Q86bq7 drosophila
P01879	oryctolagus	P01879	2	299	104	6.2	803	104	6.2	P01879	P01879 oryctolagus
Q54947	rattus norv	Q54947	2	307	104	6.2	804	104	6.2	Q54947	Q54947 rattus norv
Q9nr44	homo sapien	Q9nr44	2	334	104	6.2	805	104	6.2	Q9NR44	Q9nr44 homo sapien
Q8byp1	mus musculus	Q8byp1	2	404	104	6.2	806	104	6.2	Q8BYP1	Q8byp1 mus musculus
Q82556	mus musculus	Q82556	2	524	104	6.2	807	104	6.2	Q82556	Q82556 mus musculus
Q921k7	mus musculus	Q921k7	2	524	104	6.2	808	104	6.2	Q921K7	Q921k7 mus musculus
Q8n7w7	homo sapien	Q8n7w7	2	600	104	6.2	809	104	6.2	Q8N7W7	Q8n7w7 homo sapien
Q61gl3	drosophila	Q61gl3	2	620	104	6.2	810	104	6.2	Q61GL3	Q61gl3 drosophila
Q8awt7	drosophila	Q8awt7	2	694	104	6.2	811	104	6.2	Q8AWT7	Q8awt7 drosophila
Q96p29	homo sapien	Q96p29	2	740	104	6.2	812	104	6.2	Q96P29	Q96p29 homo sapien
Q8x936	rattus norv	Q8x936	2	789	104	6.2	813	104	6.2	Q8X936	Q8x936 rattus norv
Q7tsi8	mus musculus	Q7tsi8	2	800	104	6.2	814	104	6.2	Q7TSI8	Q7tsi8 mus musculus
Q61851	mus musculus	Q61851	2	801	104	6.2	815	104	6.2	Q61851	Q61851 mus musculus
Q61006	mus musculus	Q61006	2	868	104	6.2	816	104	6.2	Q61006	Q61006 mus musculus
Q9w4y6	drosophila	Q9w4y6	2	998	104	6.2	817	104	6.2	Q9W4Y6	Q9w4y6 drosophila
Q7tcr8	mus musculus	Q7tcr8	2	1099	104	6.2	818	104	6.2	Q7TCR8	Q7tcr8 mus musculus
Q8t4l8	drosophila	Q8t4l8	2	1503	104	6.2	819	104	6.2	Q8T4L8	Q8t4l8 drosophila
Q8cfj3	mus musculus	Q8cfj3	2	147	103.5	6.1	820	103.5	6.1	Q8CFJ3	Q8cfj3 mus musculus
Q9bdn9	papio anubi	Q9bdn9	2	275	103.5	6.1	821	103.5	6.1	Q9BDN9	Q9bdn9 papio anubi
Q8uwl1	ictalurus p	Q8uwl1	2	280	103.5	6.1	822	103.5	6.1	Q8UWL1	Q8uwl1 ictalurus p
Q9cw80	mus sp. . f	Q9cw80	2	356	103.5	6.1	823	103.5	6.1	Q9CW80	Q9cw80 mus sp. . f
Q9qp5	mus musculus	Q9qp5	2	376	103.5	6.1	824	103.5	6.1	Q9QP5	Q9qp5 mus musculus
Q9qw78	mus sp. . f	Q9qw78	2	388	103.5	6.1	825	103.5	6.1	Q9QW78	Q9qw78 mus sp. . f
Q8nc34	homo sapien	Q8nc34	2	400	103.5	6.1	826	103.5	6.1	Q8NC34	Q8nc34 homo sapien
Q699p0	anthraea p	Q699p0	2	408	103.5	6.1	827	103.5	6.1	Q699P0	Q699p0 anthraea p
Q8bn18	homo sapien	Q8bn18	2	418	103.5	6.1	828	103.5	6.1	Q8BN18	Q8bn18 homo sapien
Q96cj3	homo sapien	Q96cj3	2	421	103.5	6.1	829	103.5	6.1	Q96CJ3	Q96cj3 homo sapien
P50895	homo sapien	P50895	2	428	103.5	6.1	830	103.5	6.1	P50895	Q50895 homo sapien
Q86vc7	homo sapien	Q86vc7	2	438	103.5	6.1	831	103.5	6.1	Q8VC7	Q86vc7 homo sapien
Q93383	xenopus lae	Q93383	2	447	103.5	6.1	832	103.5	6.1	Q93383	Q93383 xenopus lae
Q8c3v5	mus musculus	Q8c3v5	2	453	103.5	6.1	833	103.5	6.1	Q8C3V5	Q8c3v5 mus musculus
Q8ci8	mus musculus	Q8ci8	2	458	103.5	6.1	834	103.5	6.1	Q8CI8	Q8ci8 mus musculus

835	103.5	6.1	819	1	FR1_CHICK	P21804	gallus gall	908	101.5	6.0	1194	2	Q7TPV3	Q7cpv3	mus musculus
836	103.5	6.1	824	2	Q91286	Q91286	pleurodeles	909	101.5	6.0	1214	2	Q6ZQA6	Q6zqa6	mus musculus
837	103.5	6.1	940	2	Q8NFA7	Q8nfa7	homo sapien	910	101.5	6.0	1250	2	Q8BV01	Q8bv01	mus musculus
838	103.5	6.1	998	1	Q95R27	Q95r27	drosophila	911	101.5	6.0	1260	1	CAML_MOUSE	P11627	mus musculus
839	103.5	6.1	1356	1	VGR2_HUMAN	P35968	homo sapien	912	101	6.0	298	2	Q9GL74	Q9gl74	cercopithec
840	103.5	6.1	1513	2	Q9OZ70	Q9oz70	brachydanio	913	101	6.0	300	2	Q8BEX0	Q8bek0	cowpox viru
841	103.5	6.1	1612	1	ROB1_MOUSE	Q90270	mus musculus	914	101	6.0	507	2	Q96K30	Q96k30	homo sapien
842	103.5	6.1	1651	1	ROB1_HUMAN	Q89026	mus musculus	915	101	6.0	584	2	Q90989	Q90989	gallus gall
843	103.5	6.1	7105	2	Q7PXC9	Q7px9	anopheles g	916	101	6.0	600	2	Q8TBU0	Q8tbu0	homo sapien
844	103	6.1	229	2	Q7PUC4	Q7puc4	anopheles g	917	101	6.0	620	1	SMP_COTUA	Q92154	coturnix co
845	103	6.1	415	2	Q6O977	Q6o977	mus musculus	918	101	6.0	626	2	Q90880	Q90880	gallus gall
846	103	6.1	498	2	Q8BRT6	Q8brt6	mus musculus	919	101	6.0	673	2	Q6MZM2	Q6mzw2	homo sapien
847	103	6.1	606	2	Q8IRH8	Q8irh8	rattus norv	920	101	6.0	693	2	Q9UPU1	Q9upul	homo sapien
848	103	6.1	708	1	KIR2_HUMAN	Q6uwl6	homo sapien	921	101	6.0	700	1	KIR2_MOUSE	Q7tsu7	mus musculus
849	103	6.1	763	2	Q95YM9	Q95ym9	halocynthia	922	101	6.0	701	2	Q670J1	Q670j1	homo sapien
850	103	6.1	873	1	FAS2_DROME	P34082	drosophila	923	101	6.0	743	2	Q6P1M7	Q6plm7	homo sapien
851	103	6.1	986	1	Q6DGL7	Q6dgl7	brachydanio	924	101	6.0	789	1	KIR1_MOUSE	Q80w68	mus musculus
852	103	6.1	1388	2	Q7QKD0	Q7qkd0	anopheles g	925	101	6.0	796	2	Q91287	Q91287	pleurodeles
853	103	6.1	1470	1	ROB2_MOUSE	Q7cpd3	mus musculus	926	101	6.0	890	1	TYO3_HUMAN	Q66418	homo sapien
854	103	6.1	2693	2	Q8ISF3	Q8isf3	caenorhabdi	927	101	6.0	890	2	Q86VR3	Q86vr3	homo sapien
855	103	6.1	2708	2	Q8ISF4	Q8isf4	caenorhabdi	928	101	6.0	994	1	MERK_RAT	P57097	rattus norv
856	103	6.1	18519	2	Q8ISF6	Q8isf6	caenorhabdi	929	101	6.0	1214	2	Q75054	Q75054	homo sapien
857	103	6.1	18534	2	Q8ISF7	Q8isf7	caenorhabdi	930	101	6.0	1742	2	Q24463	Q24463	drosophila
858	102.5	6.1	224	2	Q86GV2	Q86gv2	xenopus lae	931	101	6.0	6048	2	Q7JN85	Q7jns5	caenorhabdi
859	102.5	6.1	318	2	Q8BE16	Q8be16	variola vir	932	101	6.0	6839	2	Q23550	Q23550	caenorhabdi
860	102.5	6.1	318	2	Q8BE16	Q8be16	variola vir	932	101	6.0	7158	2	Q23551	Q23551	caenorhabdi
861	102.5	6.1	354	1	HPL1_RAT	Q9qng4	variola min	933	101	6.0	890	2	Q8UWL1	Q8uwl1	ictalurus p
862	102.5	6.1	356	2	Q819K2	P03994	rattus norv	934	100.5	6.0	280	2	Q9BDM9	Q9bdm9	macaca neme
863	102.5	6.1	399	2	Q7QCH7	Q7qch7	branchiosto	935	100.5	6.0	323	2	Q9BDM9	Q9bdm9	macaca neme
864	102.5	6.1	413	2	Q27418	Q27418	manduca sex	936	100.5	6.0	333	2	Q90Z41	Q90z41	gallus gall
865	102.5	6.1	662	2	Q6O926	Q6o926	homo sapien	937	100.5	6.0	337	2	Q8GLZ7	Q8glz7	xenopus lae
866	102.5	6.1	992	2	Q9COL5	Q9c015	homo sapien	938	100.5	6.0	355	1	HPL1_CHICK	P07354	gallus gall
867	102.5	6.1	1176	1	KMLS_BOVIN	Q28824	bos taurus	939	100.5	6.0	370	2	Q800Y8	Q800y8	brachydanio
868	102.5	6.1	1184	2	Q81Y15	Q81y15	homo sapien	940	100.5	6.0	413	1	HEMO_HYACE	P25033	hyalophora
869	102.5	6.1	1651	1	ROB1_RAT	O55005	rattus norv	941	100.5	6.0	455	2	Q9UIR0	Q9uir0	homo sapien
870	102	6.0	305	2	Q8ZS95	Q8zs95	homo sapien	942	100.5	6.0	606	2	Q9ESS7	Q9ess7	mus musculus
871	102	6.0	314	2	Q8BEK8	Q8bek8	vaccinia vi	943	100.5	6.0	648	2	Q9EPF1	Q9epf1	mus musculus
872	102	6.0	319	2	Q80477	O80477	homo sapien	944	100.5	6.0	700	1	MEPB_HUMAN	Q16820	homo sapien
873	102	6.0	332	2	Q8TA95	O8ta95	homo sapien	945	100.5	6.0	802	1	FGRA_HUMAN	P22455	homo sapien
874	102	6.0	332	2	Q8BU81	Q8bu81	homo sapien	946	100.5	6.0	814	2	Q8EC41	Q8ec41	shewanella
875	102	6.0	334	2	Q76PA0	Q76pa0	homo sapien	947	100.5	6.0	847	1	CD22_HUMAN	P20273	homo sapien
876	102	6.0	350	2	Q8MS24	Q8ma24	drosophila	948	100.5	6.0	906	2	Q8TPY9	Q8tpy9	methanobarc
877	102	6.0	357	2	O15338	O15338	homo sapien	949	100.5	6.0	1250	2	Q88971	Q88971	mus musculus
878	102	6.0	359	2	P78410	P78410	homo sapien	950	100	5.9	307	2	Q94431	Q94431	ciona intes
879	102	6.0	454	2	Q68D85	Q68d85	homo sapien	951	100	5.9	314	2	Q8BEK7	Q8bek7	vaccinia vi
880	102	6.0	504	2	Q6GNA1	Q6nna1	drosophila	952	100	5.9	322	1	ICOL_MOUSE	Q9jhj8	mus musculus
881	102	6.0	508	2	Q9VGD2	Q9vgd2	drosophila	953	100	5.9	354	1	HPL1_HUMAN	P10915	homo sapien
882	102	6.0	626	1	MAG_MOUSE	P20917	mus musculus	954	100	5.9	404	2	Q8BLQ9	Q8blq9	mus musculus
883	102	6.0	664	2	Q9VGD3	Q9vgd3	drosophila	955	100	5.9	458	1	CD4_FANTR	P16004	pan troglod
884	102	6.0	702	1	CEA5_HUMAN	P06731	homo sapien	956	100	5.9	504	2	Q98923	Q98923	gallus gall
885	102	6.0	702	1	CEA5_HUMAN	Q8n4d0	homo sapien	957	100	5.9	584	2	Q98921	Q98921	gallus gall
886	102	6.0	757	1	KIR1_HUMAN	Q96184	homo sapien	958	100	5.9	592	2	Q9JLN5	Q9jln5	mus musculus
887	102	6.0	800	2	Q99052	Q99052	mus musculus	959	100	5.9	626	2	Q98922	Q98922	gallus gall
888	102	6.0	800	2	Q99052	Q99052	mus musculus	960	100	5.9	684	2	Q7SEGI	Q7segi	neuropora
889	102	6.0	1030	2	Q8NFA8	Q8nfa8	homo sapien	961	100	5.9	784	2	Q81063	Q81063	drosophila
890	102	6.0	1032	2	Q8AXZ4	Q8axz4	brachydanio	962	100	5.9	807	2	Q6DD66	Q6dd66	xenopus lae
891	102	6.0	1098	2	Q961D6	Q961d6	drosophila	963	100	5.9	818	1	TRKB_CHICK	Q91987	gallus gall
892	102	6.0	1249	1	Q7TMZ9	Q7tmz9	rattus norv	964	100	5.9	818	2	Q9PSV9	Q9psv9	xenopus lae
893	102	6.0	1271	1	MYPC_CHICK	Q90688	gallus gall	965	100	5.9	842	1	UNSA_HUMAN	Q6zn44	homo sapien
894	102	6.0	1332	2	Q9BN17	Q9bn17	drosophila	966	100	5.9	846	2	O57577	Q57577	cynops pyrr
895	102	6.0	1332	2	Q9VQW7	Q9vqw7	drosophila	967	100	5.9	1026	2	Q94780	Q94780	homo sapien
896	101.5	6.0	1426	2	Q9QZ11	Q9qz11	enterococcu	968	100	5.9	1100	2	Q94779	Q94779	homo sapien
897	101.5	6.0	336	1	C226_MACMU	O18906	macaca mula	969	100	5.9	1180	2	O57576	Q57576	cynops pyrr
898	101.5	6.0	413	2	Q26438	Q26438	hyalophora	970	100	5.9	1180	2	Q8TI57	Q8ti57	methanobarc
899	101.5	6.0	417	1	PVR_HUMAN	P15151	homo sapien	971	99.5	5.9	295	2	Q9GL75	Q9gl75	bos taurus
900	101.5	6.0	606	2	Q9VMN6	Q9vmn6	drosophila	972	99.5	5.9	323	2	Q9BDM4	Q9bdm4	macaca mula
901	101.5	6.0	624	2	Q9ESS6	Q9ess6	rattus norv	973	99.5	5.9	416	2	Q96360	Q96360	hyphantria
902	101.5	6.0	639	2	Q96P30	Q96p30	homo sapien	974	99.5	5.9	432	2	Q6DD87	Q6dde7	xenopus lae
903	101.5	6.0	686	1	SILB_HUMAN	Q96rl6	homo sapien	975	99.5	5.9	459	1	CD4_RABIT	P46630	oryctolagus
904	101.5	6.0	802	2	Q8TDA0	Q8tda0	homo sapien	976	99.5	5.9	549	2	Q6PFC5	Q6pfc5	mus musculus
905	101.5	6.0	829	2	Q9PSV8	Q9psv8	homo sapien	977	99.5	5.9	626	1	MAG_HUMAN	P20916	homo sapien
906	101.5	6.0	898	1	FAS2_SCHAM	Q9psv8	xenopus lae	978	99.5	5.9	876	2	Q7PW78	Q7pw78	anopheles g
907	101.5	6.0	912	1	ICA5_RABIT	P22648	schistocerc	979	99.5	5.9	1009	2	Q93250	Q93250	xenopus lae
			1184	2	Q66K08	Q66k08	mus musculus	980	99.5	5.9	1019	2	Q8BJK6	Q8bjk6	mus musculus

981	99.5	5.9	1028	2	Q62682	Q62682 rattus norv	1054	97.5	5.8	261	2	Q9W6V1	Q9W6V1 gallus gall
982	99.5	5.9	1031	2	Q80YN7	Q80YN7 mus musculu	1055	97.5	5.8	262	2	Q80T70	Q80T70 mus musculu
983	98.5	5.9	1256	2	Q9JIX1	Q9JIX1 mus musculu	1056	97.5	5.8	275	2	Q640S5	Q640S5 xenopus tro
984	99.5	5.9	1256	2	Q9JIX1	P79701 coturnix co	1057	97.5	5.8	313	2	Q90Z91	Q90Z91 brachydanio
985	99.5	5.9	1450	2	Q7QCP2	Q7QCP2 anopheles g	1058	97.5	5.8	316	2	Q8BEK1	Q8BEK1 cowpox viru
986	99.5	5.9	1569	2	Q6PAC0	Q6PAC0 mus musculu	1059	97.5	5.8	354	1	HPL1_BOVIN	P55252 bos taurus
987	99.5	5.9	1666	1	MYM1_MOUSE	Q62234 mus musculu	1060	97.5	5.8	354	1	HPL1_PIG	P10859 sus scrofa
988	99	5.9	234	2	Q8TZO9	Q8TZO9 mus sapien	1061	97.5	5.8	418	2	Q9EPN5	Q9EPN5 hydrochoeru
989	99	5.9	275	1	V055_FOWPV	P21975 fowlpox vir	1062	97.5	5.8	423	2	Q9WRU4	Q9WRU4 macaca mula
990	99	5.9	275	2	Q70H96	Q70H96 fowlpox vir	1063	97.5	5.8	457	1	CD4_SAISC	Q29037 saimiri sci
991	99	5.9	313	2	Q89197	Q89197 variola vir	1064	97.5	5.8	457	2	Q8HZT7	Q8HZT7 saimiri sci
992	99	5.9	359	1	HPL3_MOUSE	Q80WM5 mus musculu	1065	97.5	5.8	473	2	Q83HS2	Q83HS2 tropheryma
993	99	5.9	458	1	CD4_HUMAN	P01730 homo sapien	1066	97.5	5.8	473	2	Q83MX5	Q83MX5 tropheryma
994	99	5.9	520	2	Q7Z3M6	Q7Z3M6 homo sapien	1067	97.5	5.8	476	2	Q80WU0	Q80WU0 mus musculu
995	99	5.9	542	2	Q8NHN5	Q8NHN5 homo sapien	1068	97.5	5.8	510	2	Q6EH12	Q6EH12 rattus norv
996	99	5.9	584	2	Q00478	Q00478 homo sapien	1069	97.5	5.8	534	2	Q86SE4	Q86SE4 homo sapien
997	99	5.9	734	2	Q96LA4	Q96LA4 homo sapien	1070	97.5	5.8	534	2	Q866T2	Q866T2 pan troglod
998	99	5.9	734	2	Q96P31	Q96P31 homo sapien	1071	97.5	5.8	764	1	ICCR_DROME	Q08180 drosophila
999	99	5.9	742	2	Q8N6S2	Q8N6S2 homo sapien	1072	97.5	5.8	764	2	Q8MQQ1	Q8MQQ1 drosophila
1000	99	5.9	756	2	Q800Z0	Q800Z0 brachydanio	1073	97.5	5.8	764	2	Q9W4U1	Q9W4U1 drosophila
1001	99	5.9	773	2	Q8NSW7	Q8NSW7 homo sapien	1074	97.5	5.8	821	1	TRKB_MOUSE	P15209 mus musculu
1002	99	5.9	812	1	FGRI_XENLA	P22182 xenopus lae	1075	97.5	5.8	821	2	Q78E87	Q78E87 mus musculu
1003	99	5.9	812	1	Q91742	Q91742 xenopus lae	1076	97.5	5.8	876	2	Q78E87	Q78E87 mus musculu
1004	99	5.9	831	2	Q71SY9	Q71SY9 gallus gall	1077	97.5	5.8	931	1	UN5C_HUMAN	Q95185 homo sapien
1005	99	5.9	862	1	CD22_MOUSE	P35329 mus musculu	1078	97.5	5.8	1254	2	Q674V1	Q674V1 podocoryne
1006	99	5.9	937	2	Q7PX00	Q7PX00 anopheles g	1079	97.5	5.8	1348	2	Q8YA37	Q8YA37 listeria mo
1007	99	5.9	975	2	Q94537	Q94537 drosophila	1080	97	5.7	1443	1	NEO1_CHICK	Q90610 gallus gall
1008	99	5.9	1526	2	Q94538	Q94538 drosophila	1081	97	5.7	240	2	Q8WUK3	Q8WUK3 homo sapien
1009	99	5.9	3197	2	Q9WUD5	Q9WUD5 drosophila	1082	97	5.7	279	2	Q7TSF0	Q7TSF0 mus musculu
1010	98.5	5.8	279	2	Q9WUD5	Q9WUD5 homo sapien	1083	97	5.7	280	2	Q8UWL2	Q8UWL2 ictalurus p
1011	98.5	5.8	283	2	Q7QGT4	Q7QGT4 anopheles g	1084	97	5.7	304	2	Q9BE26	Q9BE26 macaca fasc
1012	98.5	5.8	285	2	Q7ZVY30	Q7ZVY30 xenopus lae	1085	97	5.7	305	2	Q7PR66	Q7PR66 anopheles g
1013	98.5	5.8	288	2	Q22385	Q22385 caenorhabdi	1086	97	5.7	307	2	Q66607	Q66607 equid herpe
1014	98.5	5.8	309	2	Q9TNS7	Q9TNS7 vaccinia vi	1087	97	5.7	310	2	Q8BE18	Q8BE18 vaccinia vi
1015	98.5	5.8	316	2	Q7Q0P9	Q7Q0P9 anopheles g	1088	97	5.7	317	2	Q8BEK2	Q8BEK2 cowpox viru
1016	98.5	5.8	327	2	Q6PCB8	Q6PCB8 homo sapien	1089	97	5.7	393	2	Q8HXR8	Q8HXR8 cercopithec
1017	98.5	5.8	413	1	HEMO_MANSE	P31398 manduca sex	1090	97	5.7	456	2	Q7PUM9	Q7PUM9 anopheles g
1018	98.5	5.8	482	2	Q741A5	Q741A5 mycobacteri	1091	97	5.7	495	1	A1BG_HUMAN	P04217 homo sapien
1019	98.5	5.8	493	2	Q6P5Y4	Q6P5Y4 homo sapien	1092	97	5.7	697	2	Q8TC35	Q8TC35 homo sapien
1020	98.5	5.8	525	2	Q7QJK5	Q7QJK5 anopheles g	1093	97	5.7	703	2	Q65XY2	Q65XY2 caenorhabdi
1021	98.5	5.8	548	2	Q9Z133	Q9Z133 rattus norv	1094	97	5.7	724	2	Q8MXU5	Q8MXU5 caenorhabdi
1022	98.5	5.8	602	1	NRG1_CHICK	Q05199 gallus gall	1095	97	5.7	739	1	VCAL_HUMAN	P19320 homo sapien
1023	98.5	5.8	611	2	Q8CVC7	Q8CVC7 streptococc	1096	97	5.7	751	2	C39294	Q39294 equid herpe
1024	98.5	5.8	648	2	Q8R2Y2	Q8R2Y2 mus musculu	1097	97	5.7	824	2	Q90749	Q90749 gallus gall
1025	98.5	5.8	1005	2	P79921	P79921 xenopus lae	1098	97	5.7	931	2	Q8NPF26	Q8NPF26 homo sapien
1026	98.5	5.8	1014	2	Q8NFA6	Q8NFA6 homo sapien	1099	97	5.7	949	1	MDCL_RAT	P60756 rattus norv
1027	98.5	5.8	1021	2	P79757	P79757 gallus gall	1100	97	5.7	1028	2	Q07409	Q07409 mus musculu
1028	98.5	5.8	1333	1	VGR1_MOUSE	P79757 gallus gall	1101	97	5.7	1147	1	KMLS_RABIT	P29294 cryocolagus
1029	98	5.8	299	2	Q70863	Q70863 anopheles g	1102	97	5.7	1273	2	Q6R2F7	Q6R2F7 homo sapien
1030	98	5.8	347	2	Q8HXR7	Q8HXR7 homo sapien	1103	97	5.7	1274	1	MYPC_HUMAN	Q14896 homo sapien
1031	98	5.8	394	2	Q8HXR7	Q8HXR7 macaca fasc	1104	97	5.7	1274	2	Q9UM53	Q9UM53 homo sapien
1032	98	5.8	394	2	Q8HXR9	Q8HXR9 macaca mula	1105	97	5.7	1345	2	Q8VCD0	Q8VCD0 mus musculu
1033	98	5.8	413	2	Q8BGB4	Q8BGB4 m mus muscu	1106	97	5.7	1367	1	VGR2_MOUSE	P35918 mus musculu
1034	98	5.8	443	2	Q8N2F4	Q8N2F4 homo sapien	1107	97	5.7	1535	2	Q23991	Q23991 drosophila
1035	98	5.8	500	2	Q6UX41	Q6UX41 homo sapien	1108	97	5.7	1860	2	Q7PQF4	Q7PQF4 anopheles g
1036	98	5.8	589	2	Q8R366	Q8R366 mus musculu	1109	96.5	5.7	6620	2	Q96AA2	Q96AA2 homo sapien
1037	98	5.8	640	2	Q8BGH8	Q8BGH8 m mus muscu	1110	96.5	5.7	187	2	Q8K4E4	Q8K4E4 mus musculu
1038	98	5.8	640	2	Q8C031	Q8C031 mus musculu	1111	96.5	5.7	193	2	Q7QGT6	Q7QGT6 anopheles g
1039	98	5.8	697	1	SILA_HUMAN	Q961C7 homo sapien	1112	96.5	5.7	313	2	Q8AYZ7	Q8AYZ7 variola vir
1040	98	5.8	847	2	Q8N475	Q8N475 homo sapien	1113	96.5	5.7	313	2	Q9W6V2	Q9W6V2 gallus gall
1041	98	5.8	850	2	Q9ULF7	Q9ULF7 homo sapien	1114	96.5	5.7	352	2	Q9UQF5	Q9UQF5 homo sapien
1042	98	5.8	880	1	TYO3_MOUSE	P55144 mus musculu	1115	96.5	5.7	385	2	Q9Y3B9	Q9Y3B9 homo sapien
1043	98	5.8	880	2	Q6NZM6	P55146 rattus norv	1116	96.5	5.7	404	2	Q9J2M8	Q9J2M8 rhesus monk
1044	98	5.8	880	2	Q6NZM6	Q6NZM6 mus musculu	1117	96.5	5.7	423	2	Q5AF32	Q5AF32 leifsonia x
1045	98	5.8	931	1	UN5C_CHICK	Q7T225 gallus gall	1118	96.5	5.7	472	2	Q6AF32	Q6AF32 leifsonia x
1046	98	5.8	931	1	Q9VH85	Q9VH85 drosophila	1119	96.5	5.7	510	2	Q7L3B0	Q7L3B0 homo sapien
1047	98	5.8	945	2	Q77589	Q77589 equus cabal	1120	96.5	5.7	571	2	Q8SPI8	Q8SPI8 sus scrofa
1048	98	5.8	1021	2	Q93033	Q93033 homo sapien	1121	96.5	5.7	732	2	Q8SQ83	Q8SQ83 trichosurus
1049	98	5.8	1252	2	Q9JLI1	Q9JLI1 mus musculu	1122	96.5	5.7	772	2	Q9Y2J6	Q9Y2J6 homo sapien
1050	98	5.8	1496	2	Q90815	Q90815 gallus gall	1123	96.5	5.7	778	1	KIR3_MOUSE	Q92J6 mus musculu
1051	98	5.8	1896	2	Q9IAJ1	Q9IAJ1 xenopus lae	1124	96.5	5.7	782	2	Q9ESAS	Q9ESAS rattus norv
1052	97.5	5.8	210	2	Q7EVL9	Q7EVL9 anopheles g	1125	96.5	5.7	800	2	Q86LF9	Q86LF9 drosophila
1053	97.5	5.8	257	2	Q6NW92	Q6NW92 brachydanio	1126	96.5	5.7	801	2	Q86LF8	Q86LF8 drosophila
										804	2	Q800Z1	Q800Z1 brachydanio

1127	96.5	5.7	806	2	Q90200	Q90200 brachydanio	1200	95	5.6	879	1	FPRP_RAT	Q62786 rattus norv
1128	96.5	5.7	1200	1	HYAL_STRPU	O76536 strongyloce	1201	95	5.6	949	1	MDC1_MOUSE	P60755 mus musculu
1129	96.5	5.7	1232	2	O8TCG8	O8tcg8 homo sapien	1202	95	5.6	994	1	MERK_MOUSE	O60805 mus musculu
1130	96.5	5.7	1560	1	TENN_MOUSE	O80271 mus musculu	1203	95	5.6	1052	2	O7PMY4	O7pmv4 anopheles g
1131	96.5	5.7	1944	2	O695L3	O695l3 brachydanio	1204	95	5.6	1270	1	MYPC_MOUSE	O70468 mus musculu
1132	96.5	5.7	2013	2	O8Y9T8	O8y9t8 listeria mo	1205	95	5.6	1104	2	O9VBE5	O9vbe5 drosophila
1133	96.5	5.7	3198	2	O8Y9G8	O8y9g8 manduca sex	1206	95	5.6	1371	2	O8BV57	O8bv57 mus musculu
1134	96	5.7	228	2	O7PVK5	O7pvk5 anopheles g	1207	95	5.6	1375	2	O8ML47	O8ml47 drosophila
1135	96	5.7	234	2	O9MCS7	O9mc87 bacterioph	1208	95	5.6	1450	1	MPSP_CHICK	O02173 gallus gall
1136	96	5.7	236	2	O6P5S3	O6p5s3 homo sapien	1209	95	5.6	1501	2	O7TTI7	O7tti7 mus musculu
1137	96	5.7	252	2	O8ML12	O8ml12 drosophila	1210	95	5.6	1501	2	O9QW00	O9qw00 rattus sp.
1138	96	5.7	310	2	O7QK35	O7qk35 anopheles g	1211	95	5.6	1501	2	O9V6D5	O9v6d5 drosophila
1139	96	5.7	312	2	O8BEI7	O8bei7 vaccinia vi	1212	95	5.6	1526	2	O9VZZ4	O9vzz4 drosophila
1140	96	5.7	320	2	O8BEJ2	O8bej2 rabbitpox v	1213	95	5.6	1863	2	O64605	O64605 rattus norv
1141	96	5.7	360	2	O8N732	O8n732 homo sapien	1214	95	5.6	1898	2	O9EQ17	O9eq17 mus musculu
1142	96	5.7	395	2	O6IE78	O6ie78 paralichthy	1215	95	5.6	1904	2	O64699	O64699 mus musculu
1143	96	5.7	407	1	GLC2_VIBCH	O9klp4 vibrio chol	1216	95	5.6	1945	2	O96681	O96681 drosophila
1144	96	5.7	462	2	O7RTW1	O7rtw1 homo sapien	1217	95	5.6	22152	2	O8WXI7	O8wx17 homo sapien
1145	96	5.7	492	2	O7QD44	O7qd44 anopheles g	1218	94.5	5.6	315	1	HEMA_VACCC	P20378 vaccinia vi
1146	96	5.7	570	2	O6GMZ9	O6gmz9 xenopus lae	1219	94.5	5.6	330	2	O8C2J8	O8c2j8 mus musculu
1147	96	5.7	619	2	O7PX10	O7px10 anopheles g	1220	94.5	5.6	341	2	O8I9K3	O8i9k3 branchiosto
1148	96	5.7	636	2	O7LON3	O7lon3 homo sapien	1221	94.5	5.6	371	2	O6DCH5	O6dch5 xenopus lae
1149	96	5.7	639	1	NRG1_HUMAN	O02297 h pro-neure	1222	94.5	5.6	381	2	O9Y4A4	O9y4a4 homo sapien
1150	96	5.7	640	2	O7RTV8	O7rtv8 homo sapien	1223	94.5	5.6	388	2	O8NFZ8	O8nfz8 homo sapien
1151	96	5.7	640	2	O9HCJ2	O9hcj2 homo sapien	1224	94.5	5.6	500	2	O9XZB7	O9xz27 drosophila
1152	96	5.7	650	1	LiB1_HUMAN	O8nh16 h leukocyte	1225	94.5	5.6	553	1	IRL1_HUMAN	O8wxj5 homo sapien
1153	96	5.7	739	1	VCA1_MOUSE	P29533 mus musculu	1226	94.5	5.6	556	1	O8WXJ5	O8wxj5 homo sapien
1154	96	5.7	739	2	O91X98	O91x98 mus musculu	1227	94.5	5.6	570	2	O6GLY1	O6gly1 xenopus lae
1155	96	5.7	802	2	O95M13	O95m13 bos taurus	1228	94.5	5.6	588	1	C166_CHICK	P42292 gallus gall
1156	96	5.7	1016	2	O7JFL6	O7jfl6 oryctolagus	1229	94.5	5.6	593	2	O61NM5	O6imn5 xenopus lae
1157	96	5.7	1193	2	O9BKU8	O9bk18 aplysia cal	1230	94.5	5.6	628	2	O9MZ08	O9mz08 bos taurus
1158	96	5.7	1193	2	O9VQW1	O9vqw1 drosophila	1231	94.5	5.6	733	2	O9QZM7	O9qzm7 mus musculu
1159	96	5.7	1252	2	O9EQS9	O9eqs9 mus musculu	1232	94.5	5.6	755	2	O8CCF8	O8ccf8 mus musculu
1160	96	5.7	1253	2	O9EQS8	O9eqs8 mus musculu	1233	94.5	5.6	822	1	TRKB_HUMAN	O16320 homo sapien
1161	96	5.7	1294	2	O80TB0	O80tb0 mus musculu	1234	94.5	5.6	838	2	O8WXJ7	O8wxj7 homo sapien
1162	96	5.7	1530	2	O8Y479	O8y479 listeria mo	1235	94.5	5.6	868	2	O86VF2	O86vf2 homo sapien
1163	96	5.7	1744	2	O82TW8	O8y479 enterococc	1236	94.5	5.6	879	2	O6PEB0	O6pe80 mus musculu
1164	96	5.7	6875	2	O28733	O28733 oryctolagus	1237	94.5	5.6	888	1	UFO_MOUSE	O00993 mus musculu
1165	95.5	5.7	220	2	O7PX67	O7px67 anopheles g	1238	94.5	5.6	888	2	O8QYQ3	O80yq3 mus musculu
1166	95.5	5.7	243	1	CD48_HUMAN	P09326 homo sapien	1239	94.5	5.6	891	2	O9QV74	O9qv74 mus musculu
1167	95.5	5.7	243	2	O6IAZ2	O6iaz2 homo sapien	1240	94.5	5.6	931	1	UNSC_MOUSE	O08747 mus musculu
1168	95.5	5.7	308	2	O6QVZ1	O6qvz1 aratubaba v	1241	94.5	5.6	999	1	MERK_HUMAN	O12866 homo sapien
1169	95.5	5.7	308	2	O91ZK9	O91zr9 cantagalo o	1242	94.5	5.6	1083	2	O76698	O76698 caenorhabdi
1170	95.5	5.7	416	2	O9NPP6	O9npp6 homo sapien	1243	94.5	5.6	1256	2	O35158	O35158 rattus norv
1171	95.5	5.7	611	2	O70W32	O70w32 oncorhynch	1244	94.5	5.6	1327	2	O8QHL3	O8qhl3 gallus gall
1172	95.5	5.7	669	2	O6B515	O6b515 poephila gu	1245	94	5.6	252	2	O8PE55	O86pe5 drosophila
1173	95.5	5.7	692	2	O800Y9	O800y9 brachydanio	1246	94	5.6	275	2	O7PVM4	O7pvm4 anopheles g
1174	95.5	5.7	757	2	O7QC00	O7qc00 anopheles g	1247	94	5.6	278	2	O9QYL3	O9qyl3 mus musculu
1175	95.5	5.7	879	2	O8V199	O8v199 rattus norv	1248	94	5.6	294	2	O8BH36	O8bh36 mesocricetu
1176	95.5	5.7	888	2	O8V1A0	O8v1a0 rattus norv	1249	94	5.6	313	1	HEMA_VARV	P33807 variola vir
1177	95.5	5.7	891	2	O8BRX1	P16419 gallus gall	1250	94	5.6	313	2	O35531	O35531 rattus norv
1178	95.5	5.7	1131	1	MYPF_CHICK	O94hms oryza sativ	1251	94	5.6	313	2	O8AYW3	O8ayw3 variola vir
1179	95.5	5.7	1163	2	O94HM5	O94hm5 oryza sativ	1252	94	5.6	343	2	O8BEI4	O8bei4 variola vir
1180	95.5	5.7	1163	2	O7XG19	O7xg19 oryza sativ	1253	94	5.6	362	2	O6U7R4	O6u7r4 mus musculu
1181	95.5	5.7	1264	1	NPN_MOUSE	O9qz87 mus musculu	1254	94	5.6	377	2	O8CIC7	O8cic7 mus musculu
1182	95.5	5.7	1264	2	P91767	P91767 manduca sex	1255	94	5.6	398	2	O8QV04	O8qv04 mus musculu
1183	95.5	5.7	1485	1	O6A031	O6a031 mus musculu	1256	94	5.6	398	2	O7Z3B9	O7z3b9 homo sapien
1184	95.5	5.7	1948	1	PTNS_HUMAN	O13332 homo sapien	1257	94	5.6	419	2	O86M56	O86m56 drosophila
1185	95	5.6	181	2	O91655	O91655 xenopus lae	1258	94	5.6	484	2	O93Q08	O93q08 mus musculu
1186	95	5.6	229	2	O7PVL3	O7pvl3 anopheles g	1259	94	5.6	508	2	O8BJA5	O8bja5 mus musculu
1187	95	5.6	252	2	O9CX63	O9cx63 mus musculu	1260	94	5.6	536	2	O8BJE2	O8bj2 mus musculu
1188	95	5.6	270	2	O00426	O00426 homo sapien	1261	94	5.6	583	2	O22889	O22889 arabidopsis
1189	95	5.6	312	2	O8BEJ3	O8bej3 rabbitpox v	1262	94	5.6	583	2	O8VZR7	O8vzr7 arabidopsis
1190	95	5.6	438	2	O920C3	O920c3 mus musculu	1263	94	5.6	602	2	O9VFD9	O9vfd9 drosophila
1191	95	5.6	457	2	O960D1	O960d1 drosophila	1264	94	5.6	821	2	O8BFT0	O8bft0 m mus muscu
1192	95	5.6	521	1	C166_RABIT	O46651 oryctolagus	1265	94	5.6	883	2	O8FTT9	O8ftt9 methanosarc
1193	95	5.6	591	2	O9UQL3	O9uql3 homo sapien	1266	94	5.6	887	2	O8YWI7	O8ywi7 anabaena sp
1194	95	5.6	601	2	O86X29	O86x29 homo sapien	1267	94	5.6	943	2	O30320	O30320 archaeoglob
1195	95	5.6	814	2	O81VU1	O81vu1 homo sapien	1268	94	5.6	1010	1	CONT_CHICK	P14781 gallus gall
1196	95	5.6	814	2	O91897	O91897 xenopus lae	1269	94	5.6	1280	2	O9EPX2	O9epx2 mus musculu
1197	95	5.6	847	2	O8BFR2	O8bfr2 m mus muscu	1270	94	5.6	1325	1	YDBK_ECOLI	P32051 escherichia
1198	95	5.6	847	2	O8C4T3	O8c4t3 mus musculu	1271	94	5.6	1343	1	VGK2_RAT	O08775 rattus norv
1199	95	5.6	879	1	FPRP_MOUSE	O9wv91 mus musculu	1272	94	5.6	1406	2	O9GPF7	O9gpp7 drosophila

1273	94	5-6	1463	2	Q9VQ08	Q9vc08 drosophila	1346	92	5.5	222	2	Q8IX38	Q8ix38 homo sapien
1274	94	5-6	1945	2	Q9V491	Q9v491 drosophila	1347	92	5.5	290	2	Q7OCM3	Q7qcm3 anopheles g
1275	93.5	5-5	259	2	Q8UVA5	Q8uva5 brachydanio	1348	92	5.5	293	2	Q8AXN8	Q8axn8 cyprinus ca
1276	93.5	5-5	261	2	Q8AUQ4	Q8auq4 brachydanio	1349	92	5.5	305	2	Q8XN261	Q8xnl2 homo sapien
1277	93.5	5-5	296	2	Q42404	Q42404 gallus gall	1350	92	5.5	305	2	Q8VBM0	Q8vbm0 mus musculus
1278	93.5	5-5	313	2	Q8UV33	Q8uv33 brachydanio	1351	92	5.5	321	2	Q8MY16	Q8my16 ascaris suu
1279	93.5	5-5	315	2	Q6UN41	Q6un41 cowpox viru	1352	92	5.5	327	1	CD47 HUMAN	Q8m722 homo sapien
1280	93.5	5-5	326	2	Q8UV70	Q8uv70 brachydanio	1353	92	5.5	327	2	Q8UV63	Q8uv63 brachydanio
1281	93.5	5-5	329	1	CD86 HUMAN	P42081 homo sapien	1354	92	5.5	330	1	CD22_PONPY	Q9nl83 pongo pygma
1282	93.5	5-5	379	2	Q9CWW1	Q9cww1 mus musculus	1355	92	5.5	367	2	Q6ZW14	Q6zw14 homo sapien
1283	93.5	5-5	388	2	Q8R464	Q8r464 mus musculus	1356	92	5.5	405	2	Q6PFK4	Q6pfk4 brachydanio
1284	93.5	5-5	492	2	Q9ET54	Q9et54 mus musculus	1357	92	5.5	419	2	Q68CR6	Q68cr6 homo sapien
1285	93.5	5-5	510	2	Q801V8	Q801v8 brachydanio	1358	92	5.5	422	2	Q7RTV9	Q7rtv9 homo sapien
1286	93.5	5-5	655	2	Q8A9U5	Q8a9u5 bacteroides	1359	92	5.5	485	2	Q801W5	Q801w5 brachydanio
1287	93.5	5-5	729	1	FGRL DROME	Q07407 drosophila	1360	92	5.5	526	2	Q8NG09	Q8ng09 homo sapien
1288	93.5	5-5	739	2	Q28260	Q28260 canis famli	1361	92	5.5	544	2	Q7ZZ85	Q7zz85 brachydanio
1289	93.5	5-5	778	1	KIR3_HUMAN	Q8liu9 homo sapien	1362	92	5.5	564	1	C166_BRARE	Q90460 brachydanio
1290	93.5	5-5	808	1	FGRA_MOUSE	Q03142 mus musculus	1363	92	5.5	564	2	C161QX4	Q6lqx4 brachydanio
1291	93.5	5-5	853	2	Q6DPX7	Q6dfx7 mus musculus	1364	92	5.5	583	1	C166_HUMAN	Q13740 homo sapien
1292	93.5	5-5	1040	2	Q9W675	Q9w675 brachydanio	1365	92	5.5	613	2	Q969F0	Q969p0 homo sapien
1293	93.5	5-5	1331	2	Q7Q623	Q7q623 anopheles g	1366	92	5.5	696	1	IP11_RAT	P59824 rattus norv
1294	93.5	5-5	1630	2	Q9OT24	Q9ot24 gallus gall	1367	92	5.5	700	2	Q9F244	Q9f244 homo sapien
1295	93.5	5-5	1817	2	Q8TI59	Q8ti59 methanosarc	1368	92	5.5	812	2	Q69ZJ6	Q69zj6 mus musculus
1296	93.5	5-5	5327	1	MACF_MOUSE	Q9qzx0 mus musculus	1369	92	5.5	828	2	Q91743	Q91743 xenopus lae
1297	93	5-5	280	2	Q8UV51	Q8uv51 brachydanio	1370	92	5.5	902	2	Q17576	Q17576 caenorhabdi
1298	93	5-5	302	1	ICOL_HUMAN	Q75144 homo sapien	1371	92	5.5	928	2	Q9BLX1	Q9bly1 caenorhabdi
1299	93	5-5	308	2	Q9IC17	Q9ici7 vaccinia vi	1372	92	5.5	931	1	UN5C_RAT	Q761x5 rattus norv
1300	93	5-5	333	2	Q7PXA4	Q7pxa4 anopheles g	1373	92	5.5	956	1	MDC1_HUMAN	Q74353 homo sapien
1301	93	5-5	336	2	Q8WVY5	Q8wv5 homo sapien	1374	92	5.5	3396	1	PGCV_HUMAN	P13611 homo sapien
1302	93	5-5	388	1	BASI_RAT	P26453 rattus norv	1375	92	5.5	12268	2	Q8MQ08	Q8mq08 caenorhabdi
1303	93	5-5	390	2	Q7QBR2	Q7qbr2 anopheles g	1376	92	5.5	13100	2	Q9R165	Q9r165 caenorhabdi
1304	93	5-5	413	2	Q6KDB4	Q6kdb4 escherichia	1377	91.5	5.4	153	2	Q9R1A2	Q9r1a2 mus musculus
1305	93	5-5	424	2	Q8FIX7	Q8fix7 escherichia	1378	91.5	5.4	196	2	Q7PJY5	Q7pjy5 anopheles g
1306	93	5-5	435	1	PSG6_HUMAN	Q00889 homo sapien	1379	91.5	5.4	234	2	Q6NS95	Q6ns95 homo sapien
1307	93	5-5	459	2	Q6ZMD0	Q6zmd0 homo sapien	1380	91.5	5.4	240	1	CD48_MOUSE	P18181 mus musculus
1308	93	5-5	523	2	O00480	O00480 homo sapien	1381	91.5	5.4	240	2	Q6P905	Q6p905 mus musculus
1309	93	5-5	555	1	C166_CARAU	Q90304 carassius a	1382	91.5	5.4	299	2	Q9VHQ8	Q9vhq8 drosophila
1310	93	5-5	569	2	Q8AXU1	Q8axu1 oncorhynch	1383	91.5	5.4	330	1	EMB_MOUSE	P21995 mus musculus
1311	93	5-5	606	2	Q9BZ20	Q9bz20 homo sapien	1384	91.5	5.4	330	2	Q8C543	Q8c543 mus musculus
1312	93	5-5	606	2	Q8BZD4	Q8bz4 m mus muscu	1385	91.5	5.4	366	2	Q64216	Q64216 spalax zemn
1313	93	5-5	650	2	Q9GKR2	Q9gkr2 bos taurus	1386	91.5	5.4	412	2	Q8HXT9	Q8hxt9 tolypeutes
1314	93	5-5	739	2	Q9GKR3	Q9gkr3 bos taurus	1387	91.5	5.4	451	2	Q8VDS1	Q8vds1 mus musculus
1315	93	5-5	739	2	Q8K0X1	Q8k0x1 mus musculus	1388	91.5	5.4	459	2	Q86X91	Q86x91 homo sapien
1316	93	5-5	741	2	Q8KTF0	Q8ktf0 listeria mo	1389	91.5	5.4	463	2	Q9GMB5	Q9gmb5 bos indicus
1317	93	5-5	848	2	Q25198	Q25198 hydra atten	1390	91.5	5.4	541	2	Q95XJ7	Q95xj7 caenorhabdi
1318	93	5-5	1018	2	Q28106	Q28106 bos taurus	1391	91.5	5.4	570	2	Q8NCE6	Q8nce6 homo sapien
1319	93	5-5	2738	1	PGCV_RAT	Q9erb4 rattus norv	1392	91.5	5.4	573	2	Q8CEP3	Q8cep3 mus musculus
1320	93	5-5	3381	1	PGCV_BOVIN	P81282 bos taurus	1393	91.5	5.4	576	1	IL1R_MOUSE	P13504 mus musculus
1321	92.5	5-5	243	1	CAVT_BRALA	P05548 branchiost	1394	91.5	5.4	603	2	Q8VBZ9	Q8vzb9 mus musculus
1322	92.5	5-5	245	1	MOG_RAT	Q63345 rattus norv	1395	91.5	5.4	675	2	Q7T0V5	Q7tov5 xenopus lae
1323	92.5	5-5	245	2	Q6MFX9	Q6mfx9 rattus norv	1396	91.5	5.4	732	2	Q8CAW4	Q8caw4 mus musculus
1324	92.5	5-5	281	2	Q8CUE8	Q8cj8 mesocricetu	1397	91.5	5.4	789	2	Q8N2P7	Q8nzp7 homo sapien
1325	92.5	5-5	282	2	Q727D3	Q7z7d3 homo sapien	1398	91.5	5.4	826	2	Q7Q1P7	Q7q1p7 anopheles g
1326	92.5	5-5	310	2	Q8TQ07	Q8tq07 methanosarc	1399	91.5	5.4	880	2	Q8QFP9	Q8qfp9 xenopus lae
1327	92.5	5-5	330	2	Q90242	Q90242 gallus gall	1400	91.5	5.4	1048	2	Q8K135	Q8k135 mus musculus
1328	92.5	5-5	463	2	Q8CIV9	Q8civ9 mus musculus	1401	91.5	5.4	1085	2	Q8BHU7	Q8bh7 mus musculus
1329	92.5	5-5	474	2	Q8K178	Q8k178 mus musculus	1402	91.5	5.4	1085	2	Q8BH23	Q8bh23 mus musculus
1330	92.5	5-5	492	2	Q99KT6	Q99kt6 mus musculus	1403	91.5	5.4	1129	2	Q80UX0	Q80ux0 mus musculus
1331	92.5	5-5	500	2	Q9W260	Q9w260 drosophila	1404	91.5	5.4	1241	1	NPHN_HUMAN	Q60500 homo sapien
1332	92.5	5-5	621	2	Q811T7	Q811t7 mus musculus	1405	91.5	5.4	1502	2	Q9UM81	Q9um81 homo sapien
1333	92.5	5-5	669	2	Q6NN86	Q6nn86 drosophila	1406	91.5	5.4	2013	2	Q92EK2	Q92ek2 listeria in
1334	92.5	5-5	713	2	Q90330	Q90330 coturnix co	1407	91.5	5.4	2326	1	PGG2_RAT	Q00657 rattus norv
1335	92.5	5-5	762	2	Q71TW8	Q71tw8 homo sapien	1408	91.5	5.4	3347	2	Q8MMJ9	Q8mmj9 bombyx mori
1336	92.5	5-5	763	2	Q9VII4	Q9vil4 drosophila	1409	91.5	5.4	3354	2	Q8TI01	Q8ti01 bombyx mori
1337	92.5	5-5	764	1	PIGR_HUMAN	P01833 homo sapien	1410	91.5	5.4	4983	2	O11993	O11993 bovine vira
1338	92.5	5-5	764	2	Q8IZY7	Q8izy7 homo sapien	1411	91.5	5.4	163	2	Q93418	Q93418 gallus gall
1339	92.5	5-5	804	2	Q8W447	Q8w447 homo sapien	1412	91	5.4	240	2	Q7PRJ5	Q7prj5 anopheles g
1340	92.5	5-5	886	2	Q9VM64	Q9vm64 drosophila	1413	91	5.4	277	2	Q8C6H8	Q8c6h8 mus musculus
1341	92.5	5-5	891	2	Q9UHW6	Q9uhw6 homo sapien	1414	91	5.4	281	2	P97300	P97300 mus musculus
1342	92.5	5-5	1035	2	Q9NEG1	Q9neg1 drosophila	1415	91	5.4	282	2	Q9UFM8	Q9ufm8 homo sapien
1343	92.5	5-5	1053	2	Q6EQM4	Q6eqm4 oryza sativ	1416	91	5.4	309	2	Q7QFT7	Q7qft7 anopheles g
1344	92.5	5-5	1177	2	Q95YK1	Q95yk1 ciona savig	1417	91	5.4	313	2	Q9U964	Q9u964 geodia cydo
1345	92.5	5-5	1321	2	Q75129	Q75129 homo sapien	1418	91	5.4				

1419	91	5.4	398	2	Q9Y640	Q9Y640 homo sapien
1420	91	5.4	413	2	Q640R3	Q640R3 mus musculus
1421	91	5.4	416	2	Q8N7I3	Q8N7I3 homo sapien
1422	91	5.4	426	2	Q64HX5	Q64HX5 oncorhynch
1423	91	5.4	428	2	Q6F3J3	Q6F3J3 mus musculus
1424	91	5.4	573	2	Q6GN50	Q6GN50 xenopus lae
1425	91	5.4	589	2	Q6GQU6	Q6GQU6 mus musculus
1426	91	5.4	617	1	LR21_RAT	Q9Jmh2 rattus norv
1427	91	5.4	695	1	IPL1_MOUSE	P59823 mus musculus
1428	91	5.4	736	2	Q8MYS2	Q8MYS2 drosophila
1429	91	5.4	771	1	PIGR_MOUSE	Q70570 mus musculus
1430	91	5.4	774	2	Q9V930	Q9V930 drosophila
1431	91	5.4	801	2	Q9Z0M3	Q9Z0M3 mus musculus
1432	91	5.4	824	1	MLT1_HUMAN	Q9udy8 homo sapien
1433	91	5.4	935	2	Q677R2	Q677R2 lymphocysti
1434	90.5	5.4	200	2	Q8WWT7	Q8WWT7 homo sapien
1435	90.5	5.4	226	2	Q8N440	Q8N440 homo sapien
1436	90.5	5.4	247	2	Q7PVM3	Q7PVM3 anopheles g
1437	90.5	5.4	293	2	Q7QC80	Q7QC80 anopheles g
1438	90.5	5.4	309	2	Q6YI81	Q6YI81 vaccinia vi
1439	90.5	5.4	330	2	Q96C38	Q96C38 mus musculus
1440	90.5	5.4	341	2	Q92XW0	Q92XW0 rhizobium m
1441	90.5	5.4	360	2	Q61565	Q61565 mus musculus
1442	90.5	5.4	361	2	Q9QW79	Q9QW79 mus sp. f
1443	90.5	5.4	424	1	PSGA_HUMAN	Q15235 homo sapien
1444	90.5	5.4	461	2	Q35947	Q35947 mesocricetu
1445	90.5	5.4	501	2	Q6NUV7	Q6NUV7 brachydanio
1446	90.5	5.4	584	2	Q6PGX3	Q6PGX3 brachydanio
1447	90.5	5.4	631	1	VGLF_PHODV	P28886 phocine dis
1448	90.5	5.4	655	2	Q692J1	Q692J1 streptococc
1449	90.5	5.4	655	2	Q692J6	Q692J6 streptococc
1450	90.5	5.4	655	2	Q692K1	Q692K1 streptococc
1451	90.5	5.4	662	2	Q8MJZ6	Q8MJZ6 pan troglod
1452	90.5	5.4	687	2	Q7ZTN4	Q7ZTN4 xenopus lae
1453	90.5	5.4	697	2	Q922E0	Q922E0 mus musculus
1454	90.5	5.4	711	2	Q24205	Q24205 drosophila
1455	90.5	5.4	717	2	Q7PUQ1	Q7PUQ1 anopheles g
1456	90.5	5.4	727	1	PEC1_MOUSE	Q08481 mus musculus
1457	90.5	5.4	771	2	Q8N116	Q8N116 homo sapien
1458	90.5	5.4	773	2	Q8ZXV2	Q8ZXV2 pyrobaculum
1459	90.5	5.4	773	2	Q7QBL9	Q7QBL9 anopheles g
1460	90.5	5.4	821	1	TRKB_RAT	Q63604 rattus norv
1461	90.5	5.4	868	1	NRG2_RAT	Q35569 rattus norv
1462	90.5	5.4	922	2	Q90413	Q90413 brachydanio
1463	90.5	5.4	1043	2	Q6PA07	Q6PA07 xenopus lae
1464	90.5	5.4	1205	2	Q8BUJ0	Q8BUJ0 mus musculus
1465	90.5	5.4	1282	2	Q90X86	Q90X86 xenopus lae
1466	90.5	5.4	1343	2	Q8G524	Q8G524 bifidobacte
1467	90.5	5.4	1383	2	Q7Q840	Q7Q840 anopheles g
1468	90.5	5.4	1878	2	Q6CPZ4	Q6CPZ4 kluyveromyc
1469	90.5	5.4	2046	2	Q757C8	Q757C8 ashbya gos
1470	90.5	5.4	3029	2	Q7Q767	Q7Q767 anopheles g
1471	90	5.3	96	2	Q08909	Q08909 mus musculus
1472	90	5.3	203	2	Q7PVM1	Q7PVM1 anopheles g
1473	90	5.3	249	2	Q8R477	Q8R477 mus musculus
1474	90	5.3	249	2	Q8R2R4	Q8R2R4 mus musculus
1475	90	5.3	288	2	Q28499	Q28499 macaca mula
1476	90	5.3	290	2	Q9EP73	Q9EP73 mus musculus
1477	90	5.3	304	2	Q8BID4	Q8BID4 mus musculus
1478	90	5.3	416	2	Q67IP8	Q67IP8 homo sapien
1479	90	5.3	422	2	Q8WR61	Q8WR61 lymantria d
1480	90	5.3	468	2	Q7TNR6	Q7TNR6 mus musculus
1481	90	5.3	484	2	Q8BSF0	Q8BSF0 mus musculus
1482	90	5.3	494	2	Q9ESC6	Q9ESC6 mus musculus
1483	90	5.3	510	2	Q8BUC9	Q8BUC9 mus musculus
1484	90	5.3	526	2	Q23333	Q23333 caenorhabdi
1485	90	5.3	531	2	Q7KYN4	Q7KYN4 homo sapien
1486	90	5.3	558	2	Q6NUX0	Q6NUX0 brachydanio
1487	90	5.3	571	2	Q8QFN4	Q8QFN4 salmo salar
1488	90	5.3	585	2	Q6TS33	Q6TS33 brachydanio
1489	90	5.3	593	2	Q9WU74	Q9WU74 rattus norv
1490	90	5.3	609	2	Q7QHA0	Q7QHA0 anopheles g
1491	90	5.3	618	1	LR21_MOUSE	Q8K099 mus musculus
1492	90	5.3	645	2	Q8BUJ6	Q8BUJ6 mus musculus
1493	90	5.3	659	2	Q8BIH3	Q8BIH3 mus musculus
1494	90	5.3	723	2	Q86YI4	Q86YI4 homo sapien
1495	90	5.3	737	2	Q965M3	Q965M3 caenorhabdi
1496	90	5.3	821	1	FGR2_HUMAN	P21802 homo sapien
1497	90	5.3	845	1	MCDL_HUMAN	Q9HBB8 homo sapien
1498	90	5.3	881	2	Q965M2	Q965M2 caenorhabdi
1499	90	5.3	943	1	UN5B_XENLA	Q8JGT4 xenopus lae
1500	90	5.3	1018	1	CONT_HUMAN	Q12860 homo sapien
ALIGNMENTS						
RESULT 1						
Q6UXI4						
ID	Q6UXI4	PRELIMINARY;	PRT;	321 AA.		
AC	Q6UXI4;					
DT	05-JUL-2004 (TREMBLrel. 27, Created)					
DR	05-JUL-2004 (TREMBLrel. 27, Last sequence update)					
DE	STIGMA.					
GN	ORFNames=UNQ317;					
OS	Homo sapiens (Human)					
OC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;					
OC	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.					
OX	NCBI_TaxID=9606;					
RP	[1]					
RN	SEQUENCE FROM N.A.					
RX	MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;					
RA	Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,					
RA	Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,					
RA	Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,					
RA	Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,					
RA	Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,					
RA	Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,					
RA	Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D.,					
RA	Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,					
RA	Godowski P.;					
RT	"The secreted protein discovery initiative (SPDI), a large-scale					
RT	effort to identify novel human secreted and transmembrane proteins: a					
RT	bioinformatics assessment.";					
RL	Genome Res. 13:2265-2270(2003).					
DR	EMBL; AY358341; AAQ88707.1; -.					
DR	InterPro; IPR003599; IG.					
DR	InterPro; IPR007110; IG-like.					
DR	InterPro; IPR003598; IG_C2.					
DR	InterPro; IPR003596; IG_V.					
DR	Pfam; PF00047; IG; 1.					
DR	SMART; SM00409; IGC2; 1.					
DR	SMART; SM00408; IGC2; 1.					
DR	SMART; SM00406; IGV; 1.					
DR	PROSITE; PS50835; IG LIKE; 2.					
SQ	SEQUENCE 321 AA; 35544 MW; B2AB2E3151D39C6E CRC64;					
Query Match 100.0%; Score 1688; DB 2; Length 321;						
Best Local Similarity 100.0%; Pred. No. 1.7e-125;						
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;						
Qy	1	MGILLGLLLGLLTVDTYGRFLEVPESVTGPKGDVNLPCITYDPLQGYTVLVKVLVQR	60			
Db	1	MGILLGLLLGLLTVDTYGRFLEVPESVTGPKGDVNLPCITYDPLQGYTVLVKVLVQR	60			
Qy	61	GSDPTVIFLRSSGGDHIOQAKYQGRHLVSHKVPDVSILQLSTLEMDDRSHYTCVTTWTP	120			
Db	61	GSDPTVIFLRSSGGDHIOQAKYQGRHLVSHKVPDVSILQLSTLEMDDRSHYTCVTTWTP	120			
Qy	121	DGNQVVRDKITELRVQKLSVSKPTVTTGSGYFTVPQGMRLSLQCAQSGSPPISYWKQ	180			
Db	121	DGNQVVRDKITELRVQKLSVSKPTVTTGSGYFTVPQGMRLSLQCAQSGSPPISYWKQ	180			
Qy	181	QTNNOEPIKATVLTSTLLPKPAVIAVDSGYFTAGQVSGEHSIDIVKFWKDSKLLTKK	240			

Db 181 QTNQNEPIKATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKXSSKLLKTK 240
QY 241 TEAPTTMTYPLKATSTVKQSWDWTMDGGLGETSAGPKSLPVPFAILLIISLCCMVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSWDWTMDGGLGETSAGPKSLPVPFAILLIISLCCMVFT 300
QY 301 MAYIMLCRKTSQOEHVYEAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAR 321

RESULT 2

QY279 ID Q9Y279 PRELIMINARY; PRT; 399 AA.
AC Q9Y279
DT 01-NOV-1999 (TRENBLrel. 12, Created)
DT 01-NOV-1999 (TRENBLrel. 12, Last sequence update)
DT 25-OCT-2004 (TRENBLrel. 28, Last annotation update)
DE 239Ig protein precursor (V-set and immunoglobulin domain containing 4).
DE
GN Name=239Ig; Synonyms=VSI64;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RV
SEQUENCE FROM N.A.
RX MEDLINE=20461865; PubMed=11004523; DOI=10.1016/S0167-4781(00)00131-7;
RA Langane K., Colleaux L., Kloos D.U., Fontes M., Wiesacker P.;
RT "Cloning of 239Ig, a novel gene with immunoglobulin-like domains
RT located on human chromosome X";
RL Biochim. Biophys. Acta 1492:522-525(2000).
RN [2]

SEQUENCE FROM N.A.

RC TISSUE=Brain;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Villalon D.K., Muzny D.M., Halse S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RA "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [3]

SEQUENCE FROM N.A.

RC TISSUE=Brain;
RX Director MGC Project;
RA Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.

DR EMBL; AJ132502; CAB51536.1; -;
DR EMBL; BC018525; AAH10525.1; -;
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; Ig; 1.
DR SMART; SM00408; Igc2; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
KW Signal.
FT SIGNAL 1 19 Potential.
FT CHAIN 20 399 239Ig protein.
SEQUENCE 399 AA; 43987 MW; 735CA3BC58180535 CRC64;

Query Match 100.0%; Score 1688; DB 2; Length 399;
Best Local Similarity 100.0%; Pred. No. 2,2e-125;
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MGILLGGLLGLTVDYTGRIPLVPSVTGPKWGDVNLPTCTYDPLQYTVLVKWLVR 60
Db 1 MGILLGGLLGLTVDYTGRIPLVPSVTGPKWGDVNLPTCTYDPLQYTVLVKWLVR 60
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWQTP 120
Db 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWQTP 120
QY 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYGFVPOQMRISLQCOARGSPPIISYWKQ 180
Db 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYGFVPOQMRISLQCOARGSPPIISYWKQ 180
QY 181 QTNQNEPIKATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKXSSKLLKTK 240
Db 181 QTNQNEPIKATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKXSSKLLKTK 240
QY 241 TEAPTTMTYPLKATSTVKQSWDWTMDGGLGETSAGPKSLPVPFAILLIISLCCMVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSWDWTMDGGLGETSAGPKSLPVPFAILLIISLCCMVFT 300
QY 301 MAYIMLCRKTSQOEHVYEAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAR 321

RESULT 3

Q80WA3 ID Q80WA3 PRELIMINARY; PRT; 280 AA.
AC Q80WA3
DT 01-JUN-2003 (TRENBLrel. 24, Created)
DT 01-JUN-2003 (TRENBLrel. 24, Last sequence update)
DT 01-MAR-2004 (TRENBLrel. 26, Last annotation update)
DE Hypothetical protein BC025105.
GN Name=BC025105;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RV
SEQUENCE FROM N.A.
RC STRAIN=FVB/N; TISSUE=Liver;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RA "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RV
SEQUENCE FROM N.A.
RC STRAIN=FVB/N; TISSUE=Liver;
RX Strausberg R.;
RA Submitted (MAR-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; BC025105; AAH25105.1; -;
DR MGD; MGI:2679720; BC025105.

DR	InterPro; IPR007110; Ig-like.
DR	PROSITE; PS00835; IG_LIKE; 1.
KW	Hypothetical protein.
SQ	SEQUENCE 280 AA; 31467 MW; 3F099AE996CEB977 CRC64;
 Query Match 43.1%; Score 728; DB 2; Length 280; Best Local Similarity 47.0%; Pred.No. 1.6e-49; Matches 151; Conservative 29; Mismatches 45; Indels 96; Gaps 2	
QY	1 MGILLGLLLCHLVTDVTGRPILEVPSVTPGKGDVNLPCTYPDLOGYTQVLVKWLVR 60 : :
Dd	1 MEISSGLFLGHILVLYTGHTLKTSPSVGTGWKDVIKI QIYPDLRGYRQLVKWLVRH 60 :
QY	61 GSDPVITIFLRDSSGDHIQQAKYQGRLHVSHKVPDGVSLSLTLEMDRRSHYTCVWTQP 120 : :
Dd	61 GSDSVTIFLRDSTGDHIQQAKYRKLVSKHPVGDSLSQINTLMQDDRHNHYTCVWTQP 120 : :
QY	121 DGNQVVDPKITELRVOKLSYSKVPTVTGSYGVFPPQMRLISLCQAAGSPSIVTWKY 180 : :
Dd	121 DGNQVIDDKIIELRVRY-----NPP----- 141
QY	181 QTNNQEIPKVATSLTLFPKPAVIADSGSYFCYTAKEGVSEHQSDIVKFVWDKSLLKTK 240
Dd	142 -----RIN 144
QY	241 TEAPTTMTYPLKATTVKQSNDWDTDMGYLGETSAGPGSLPFAILIISLCMVFT 300 : : : : : : : :
Dd	145 TEAPTTHLSSL EATTIMSTSSDLTTNGTGKLEETIAGSRNLPIFAIFIISLC CIVA V T 204
QY	301 MAYIMLCRTSQOEHVTEAR 321
Dd	205 IPYILFRCTRFFQOEYYGVSR 225
 RESULT 4 JAM1_HUMAN	
ID	JAM1_HUMAN STANDARD; PRT; 299 AA.
AC	Q9Y6Z4;
DT	16-OCT-2001 (Rel. 40, Created)
DD	16-OCT-2001 (Rel. 40, Last sequence update)
DT	25-JAN-2005 (Rel. 46, Last annotation update)
DE	Junctional adhesion molecule 1 precursor (JAM) (Platelet adhesion molecule 1) (PAM-1) (Platelet Fll receptor) (UNQ264/PRO301).
DN	Name=Flilr; Synonyms=JAM1, JCAM;
GC	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX	NCBI_TaxId=9606;
RN	[1]
RP	SEQUENCE FROM N.A.
RX	MEDLINE=9323940; PubMed=10395639;
RA	Ozaki H., Ishii K., Horiuchi H., Arai H., Kawamoto T., Okawa K.,
RA	Iwamatsu A., Kita T.;
RT	"Combined treatment of TNF-alpha and IFN-gamma causes redistribution of junctional adhesion molecule in human endothelial cells.";
RT	J. Immunol. 163:553-557(1999). [2]
RN	SEQUENCE FROM N.A.
RP	Sobočka M.B., Sobočki T., Rushbrook J.I., Banerjee P., Weiss C.,
RA	Kornecki E.;
RT	"Molecular cloning and sequencing of the cDNA of Flil receptor, a novel Ig superfamily member from human platelets.";
RT	Submitted (NOV-1999) to the EMBL/GenBank/DBAJ databases. [3]
RN	SEQUENCE FROM N.A.
RP	Naik U.P., Naik M.U., DeLeon P., Spychala J.;
RT	"Cloning and characterization of PAM-1, a novel platelet adhesion molecule involved in platelet activation.";
RT	Submitted (JUL-1999) to the EMBL/GenBank/DBAJ databases. [4]
RN	SEQUENCE FROM N.A.
RP	TTSUSE=Brain;
RX	MEDLINE=21154917; PubMed=11230166; DOI=10.1101/gr.154701;

Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glassl S., Ansoerge W., Boecher M., Bloeker H., Bausachs S., Blum H., Lauber J., Duesterhoeft A., Beyer A., Koehrer K., Strack N., Mewes H.-W., Ottenwaelder B., Obermaier B., Tampe J., Heubner D., Wambutt R., Korn B., Klein M., Poustka A.;
RT "Towards a catalog of human genes and proteins: sequencing and analysis of 500 novel complete protein coding human cDNAs.";
RL Genome Res. 11:422-435 (2001).
RN [5]
RP SEQUENCE FROM N.A.
RX MEDLINE=22887256; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J., Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P., Eaton D., Foster J., Grimaldi C., Gu Q., Hase P.E., Heldens S., Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J., Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J., Seehagiri S., Simmons L., Singh J., Smith V., Spinson J., Vagts A., Vandlen R., Watanabe C., Wisand D., Woods K., Xie M.-H., Yansura D., Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I., Godowski P., Gray A.;
RA "The secreted protein novel human secreted and transmembrane proteins: effort to identify novel human secreted and transmembrane proteins: a bioinformatics assessment";
RL Genome Res. 13:2285-2270 (2003).
RN [6]
RP SEQUENCE FROM N.A.
RX TISSUE=Ovary;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.W., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Scapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Ustin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullen S.J., Bosak S.A., McSwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J.J., Helton E., Kettaman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakeley R.W., Touchman J.W., Green E.D., Dickson W.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.N., Krzywinski M.I., Skalska U., Smalish D.E., Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RA "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
RN [7]
RP SEQUENCE OF 28-42.
MEDLINE=15340161; DOI=10.1110/ps.04682504;
RA Zhang Z., Henzel W.J.;
RT "Signal peptide prediction based on analysis of experimentally verified cleavage sites";
RL Protein Sci. 13:2819-2824 (2004).
CC -1- FUNCTION: Seems to plays a role in epithelial tight junction formation. Appears early in primordial forms of cell junctions and prevents PAR3. The association of the PAR3-PAR3 complex may prevent the interaction of PAR3 with JAM1, thereby preventing tight junction assembly (By similarity). Plays a role in regulating monocyte transmigration involved in integrity of epithelial barrier. Involved in platelet activation.
CC -1- SUBUNIT: Interacts with the first PDZ domain of PAR3. The association between PAR3 and PAR6B probably disrupts this interaction (By similarity).
CC -1- SUBCELLULAR LOCATION: Type I membrane protein (Potential).
CC -1- TISSUE SPECIFICITY: Localized at tight junctions of both epithelial and endothelial cells.
CC -1- SIMILARITY: Belongs to the immunoglobulin-like V-type domains.
CC -1- SIMILARITY: Contains 2 immunoglobulin-like V-type domains.
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DR EMBL; AF111713; RAD42050.1; -
 DR EMBL; AF207907; AAF22829.1; -
 DR EMBL; AF172398; AAD48877.1; -
 DR EMBL; AL136649; CAB66584.1; -
 DR EMBL; AY358896; RAO89255.1; -
 DR EMBL; BC001533; RAO01533.1; -
 DR PIR; A59406; S56749.
 DR PDB; INHQ; X-ray; A/B=25-233.
 DR GENE; HGNC:14685; FLIR.
 DR MIM; 605721; -.

DR GO; GO:0005911; C:intercellular junction; TAS.
 DR GO; GO:0006954; P:inflammatory response; TAS.
 DR InterPro; IPR007110; Ig-like.

DR PROSITE; PS50835; IG_LIKE; 2.
 KW 3D-structure; Direct protein sequencing; Glycoprotein;
 KW Immunoglobulin domain; Repeat; Signal; Tight junction; Transmembrane.

FT SIGNAL 1 27
 FT CHAIN 28 299 Junctional adhesion molecule 1.
 FT DOMAIN 28 238 Extracellular (Potential).
 FT TRANSMEM 239 259 Potential.
 FT DOMAIN 260 299 Cytoplasmic (Potential).
 FT DOMAIN 28 125 Ig-like V-type 1.
 FT DOMAIN 135 228 Ig-like V-type 2.
 FT DISULFID 50 109 Potential.
 FT DISULFID 153 212 Potential.
 FT CARBOHYD 185 185 N-linked (GlcNAc...) (Potential).
 SQ SEQUENCE 299 AA; 32593 MW; D95DE2FEA23D2851 CRC64;

Query Match 10.6%; Score 178.5; DB 1; Length 299;
 Best Local Similarity 26.5%; Pred. No. 6e-06;
 Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGILGLLLGLHGLTVDTYGRPILEVPESVTPGPKGVDNLPCITYDPLQYQVVLKVLVQR 60
 DB 17 LAITLLCSALGVSIVHVS-SEPEVRIPEN-----NPVKLSLAY-----SGFSSPRVEM---- 62
 QY 61 GSDPTIFLSDSGDHIOAKQGRHLVSHK-----VPGDVSLSQISTLEMDRSHYTCVEV 115
 DB 63 -----KFDGDTTLVLCVNNKITASYEDRVTFPLPTGITFKSVTRE--DTGYTCVK 111
 QY 116 TWQTPDGNQVVDKITELRVQKLSVSKPTVTGSGVGTVPQGMRIISLQCOAR-GSPPTS 174
 DB 112 SEEGSGSYGEVKKVLIIVL-----VPPSKPTVINPS-----SATIGNRAVLTCSEQDSPPSE 163
 QY 175 YIWKY-----QQTN-----NQEPKIVATSLTLFKPAVIADSGSYCTAKQGVGSQH 222
 DB 164 YTWFKDGIWMPNTPKSTRAFSNSSLVNLPTTGELVFDPLSLASDTGEYSCEARNGYGTPT 223
 QY 223 SDIVK 227
 DB 224 SNAVR 228

RESULT 5
 A33_HUMAN
 ID A33_HUMAN STANDARD; PRT; 319 AA.
 AC Q99795;
 DT 01-NOV-1997 (Rel. 35, Created)
 DT 01-NOV-1997 (Rel. 35, Last sequence update)
 DT 05-JUL-2004 (Rel. 44, Last annotation update)
 DE Cell surface A33 antigen precursor (Glycoprotein A33).
 GN Name=GPA33;
 OS Homo sapiens (Human).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 OK NCBI_TaxID=9606;

RN SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
 RP TISSUE=Colon carcinoma;
 RC MEDLINE=97165045; PubMed=9012807; DOI=10.1073/pnas.94.2.469;
 RA Heath J.K., White S.J., Johnstone C.N., Catmel B., Simpson R.J.,
 RA Moritz R.L., Tu G.-F., Ji H., Whitehead R.H., Groenen L.C.,
 RA Scott A.W., Ritter G., Cohen L., Welt S., Old L.J., Nice E.C.,
 RA Burgess A.W.;
 RT "The human A33 antigen is a transmembrane glycoprotein and a novel
 RT member of the immunoglobulin superfamily";
 RL Proc. Natl. Acad. Sci. U.S.A. 94:469-474(1997).
 RN [2]
 RP POST-TRANSLATIONAL MODIFICATIONS.
 RX MEDLINE=97396159; PubMed=9245713; DOI=10.1006/bbrc.1997.6966;
 RA Ritter G., Cohen L.S., Nice E.C., Catmel B., Burgess A.W.,
 RA Moritz R.L., Ji H., Heath J.K., White S.J., Welt S., Old L.J.,
 RA Simpson R.J.;
 RT "Characterization of posttranslational modifications of human A33
 RT antigen, a novel palmitoylated surface glycoprotein of human
 RT gastrointestinal epithelium"; Res. Commun. 236:682-686(1997).
 RL Biochem. Biophys. Res. Commun. 236:682-686(1997).
 CC -!- FUNCTION: May play a role in cell-cell recognition and signaling.
 CC -!- SUBCELLULAR LOCATION: Type I membrane protein.
 CC -!- TISSUE SPECIFICITY: Expressed in normal gastrointestinal
 CC epithelium and in 95% of colon cancers.
 CC -!- PTM: N-glycosylated, contains approximately 8 kDa of N-linked
 CC carbohydrate.
 CC -!- PTM: Palmitoylated.
 CC -!- SIMILARITY: Contains 1 immunoglobulin-like C2-type domain.
 CC -!- SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
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 CC between the Swiss Institute of Bioinformatics and the EMBL Outstation -
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DR EMBL; U79725; AAC50957.1; -.
 DR HSSP; O88792; 1F97.
 DR GENE; HGNC:4445; GPA33.
 DR MIM; 602171; -.
 DR GO; GO:0005887; C:integral to plasma membrane; TAS.
 DR GO; GO:0004872; F:receptor activity; TAS.
 DR InterPro; IPR007110; Ig-like.
 DR Pfam; PF00047; Ig; 2.
 DR PROSITE; PS50835; IG_LIKE; 2.
 KW Antigen; Direct protein sequencing; Glycoprotein;
 KW Immunoglobulin domain; Lipoprotein; Palmitate; Signal; Transmembrane.
 FT SIGNAL 1 21
 FT CHAIN 22 319 Cell surface A33 antigen.
 FT DOMAIN 22 235 Extracellular (Potential).
 FT TRANSMEM 236 256 Potential.
 FT DOMAIN 257 319 Cytoplasmic (Potential).
 FT DOMAIN 22 134 Ig-like V-type.
 FT DOMAIN 140 227 Ig-like C2-type.
 FT DOMAIN 258 261 Poly-Cys.
 FT DISULFID 43 117 Potential.
 FT DISULFID 146 222 Potential.
 FT DISULFID 162 211 Potential.
 FT CARBOHYD 112 112 N-linked (GlcNAc...) (Potential).
 FT CARBOHYD 200 200 N-linked (GlcNAc...) (Potential).
 FT CARBOHYD 223 223 N-linked (GlcNAc...) (Potential).
 SQ SEQUENCE 319 AA; 35632 MW; 9BFC7AAF45C2408E CRC64;

Query Match 10.5%; Score 177; DB 1; Length 319;
 Best Local Similarity 27.5%; Pred. No. 8.6e-06;
 Matches 64; Conservative 39; Mismatches 94; Indels 36; Gaps 11;

QY 4 LLGILLGLHGLTVDTYGRPILEVPESVTPGPKGVDNLPCITYDPLQYQVVLKVM--LVQR 60
 DB 8 VLWTLCAVRVTDAIS---VETPDVLRASQKSVTLPTCTVHTSTSSREGLIQWDKLLIT 64

KW Glycoprotein; Immunoglobulin domain; Repeat; Signal; Tight junction; Transmembrane.
FT SIGNAL 1 24 Potential.
FT CHAIN 25 298 Junctional adhesion molecule 1.
FT DOMAIN 25 237 Extracellular (Potential).
FT TRANSMEM 238 258 Potential.
FT DOMAIN 259 298 Cytoplasmic (Potential).
FT DOMAIN 28 124 Ig-like V-type 1.
FT DISULFID 134 227 Ig-like V-type 2.
FT DOMAIN 152 211 Potential.
FT CARBOHYD 184 194 N-linked (GlcNAc...) (Potential).
SQ SEQUENCE 298 AA; 32456 MW; 714FE1C1714769A2 CRC64;

Query Match 10.0%; Score 169.5; DB 1; Length 298;
Best Local Similarity 25.1%; Pred. No. 3.1e-05;
Matches 63; Conservative 31; Mismatches 88; Indels 69; Gaps 11;

QY 3 ILGLGLLLGLTVDYGRILEVPESVTPGWKGVNLCTDPLQGYTQVLVKVLVQRG- 61
DB 18 MILCSLALGRGAVQTY-BFVRVPEP-----NPAKLJSCSY---SGFSRPRVEMKFTHGD 67
QY 62 -----SDPVTIFLDRSSGDHIQAKYQGRHLVSHKVPDGLSLQLSTLMDRSHYTCE 114
DB 68 IRLGLVYNNKIT-----ASVENRTFS-----DTGITFHSVTRKDTGMYTOM 109
QY 115 VTWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGFVTPQGNRISLQCAR-GSPPI 173
DB 110 V5-----DEGNTYGEVTVQLIVLPSPKPTINVP-----SVTIGTRAVLTCSRDSGSPS 161
QY 174 SYIWK-----OQTWQEPKIVATLSTLFLKPAVITADSGSYFCTAKG 215
DB 162 EYKFKDGVEMPLEPKSNRAFSNSSTYTLNQK-----TGELIFDPSASDTGDFTCQAQN 215
QY 216 QVSGSEQSHDIV 226
DB 216 GYASPVKSDTV 226

RESULT 10
Q8WV3 PRELIMINARY; PRT; 365 AA.
AC Q8WV3;
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Cocksackie virus and adenovirus receptor BCAR.
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21547769; PubMed=11689979; DOI=10.1006/bbrc.2001.5851;
RA Thoenen I., Keyaerts E., Lindberg M., Van Ranst M.;
RT "Characterization of a cDNA encoding the bovine coxsackie and
RT adenovirus receptor";
RL Biochem. Biophys. Res. Commun. 288:805-808(2001).
DR EMBL; AY033651; AAKS7804.1; -.
DR FIR; JC7780; JC7780.
DR HSSP; P78310; 1KAC.
DR GO; GO:0004872; F:receptor activity; IEA.
DR SMART; SM00408; IGC2; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
KW Receptor.
SQ SEQUENCE 365 AA; 40153 MW; 36DE0BE5DCF86CF9 CRC64;

Query Match 10.0%; Score 169; DB 2; Length 365;
Best Local Similarity 24.8%; Pred. No. 4.4e-05;
Matches 85; Conservative 40; Mismatches 120; Indels 98; Gaps 17;

QY 1 MGILGLLLGLTVDYGRILEVPESVTPGWKGD-VNLPCY-----DPLQGYTQV 52

DB 1 MELLRLFLLCGVADFTGLSLI-TTPEQMIEKAGETAYLFECKFTLGPEDQGPLD----- 54
QY 53 LVKVLVQRGS---DPVTIFLDRSSGDHIQAKYQ---GRHLVSHK---VPGDVSLLQSTL 103
DB 55 -IEWLLSPADNQVDQVILLY---SGDKIYDDYQDLKGRVHFTSNDLKSGLASINVTNL 110
QY 104 EMDRSHYTCVETWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGFVTPQ-----GM 159
DB 111 QLSDIGTYQCKVKAPGVGNK-----KIQLTVLVKP-----SGIRCYVDGSEIGN 156
QY 160 RISLQCAR-GSPISYIWKYQQTNNQEP-----IKVATLSTLFLKPAVITADSGSYFCTAKG 215
DB 157 DFKLKCEPKESLFLRYEWQKLSQKLPSTSWLPEMTSPVISVKNASAEYSCTYCTVRN 216
QY 216 QVSGSEQSHDIVKVVVQSSKLLTKTBEATPTMTYPLKATSVTKQSDWTTDMDGYLGSTS 275
DB 217 RVGSDQ-----CLRLDVPFNSR-----AGTI 239
QY 276 AGPKSLPVPFAILLIISLCMVFTMAYIMLCRTKSQOEHVYE 318
DB 240 AG-----AVIGTLALVLIALIVE-----CCHKRREKYE 270

RESULT 11
Q8VC39 PRELIMINARY; PRT; 300 AA.
AC Q8VC39;
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE F11r protein (Mus musculus adult male cecum cDNA, RIKEN full-length
DE enriched library, clone:9130004G24 product:junction cell adhesion
DE molecule1, full insert sequence).
GN Name=F11r;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=mix FVB/N; TISSUE=Mammary tumor;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.2426038999;
RA Strauberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Uadin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny K.C., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahy J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=mix FVB/N; TISSUE=Mammary tumor;
RA Strauberg R.;
RL Submitted (JAN-2002) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RC STRAIN=CS7BL/6J; TISSUE=Cecum;
RX MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA Carninci P., Hayashizaki Y.;

[illegible]

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GenCore version 5.1.1.6
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OM protein - protein search, using sw model

Run on: August 26, 2005, 19:32:58 ; Search time 43 Seconds
(without alignments)
557.264 Million cell updates/sec

Title: US-10-767-374-2
Perfect score: 1688
Sequence: 1 MGILGLLLGLHGLTVDTYGR.....AYIMLCRKTSQOEHVEAAR 321

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1500 summaries

Database : Issued Patents AA:*
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5: /cgn2_6/prodata/1/aaa/PCTUS COMB.pep:*
6: /cgn2_6/prodata/1/aaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1688	100.0	321	4	US-09-254-465A-2
2	1688	100.0	321	4	US-09-953-499-2
3	1137	67.4	306	4	US-09-369-247-63
4	870	51.5	175	4	US-09-763-902B-6
5	178.5	10.6	299	3	US-09-188-930-189
6	178.5	10.6	299	3	US-09-188-930-331
7	178.5	10.6	299	3	US-09-462-270-2
8	178.5	10.6	299	4	US-09-254-465A-1
9	178.5	10.6	299	4	US-09-312-283C-189
10	178.5	10.6	299	4	US-09-312-283C-331
11	178.5	10.6	299	4	US-09-907-794A-119
12	178.5	10.6	299	4	US-09-905-125A-119
13	178.5	10.6	299	4	US-09-902-775A-119
14	178.5	10.6	299	4	US-09-397-243D-3
15	178.5	10.6	299	4	US-09-906-700-119
16	178.5	10.6	299	4	US-09-903-603A-119
17	178.5	10.6	299	4	US-09-904-920A-119
18	178.5	10.6	299	4	US-09-909-064-119
19	178.5	10.6	299	4	US-09-905-381A-119
20	178.5	10.6	299	4	US-09-906-618-119
21	178.5	10.6	299	4	US-09-953-499-1
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24	177	10.5	319	4	US-09-336-536-67
25	177	10.5	319	4	US-09-254-465A-6
26	177	10.5	319	4	US-09-953-499-6
27	176	10.4	320	4	US-09-254-465A-24
28	176	10.4	270	4	US-09-953-499-24
29	176	10.4	273	4	US-09-254-465A-26
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36	167	9.9	300	4	US-09-397-243D-12
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49	142.5	8.4	894	4	US-09-540-245A-15
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55	138.5	8.2	310	4	US-09-906-618-423
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58	138.5	8.2	310	4	US-09-905-125A-213
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60	138.5	8.2	360	4	US-09-906-700-213
61	138.5	8.2	360	4	US-09-903-603A-213
62	138.5	8.2	360	4	US-09-904-920A-213
63	138	8.2	360	4	US-09-905-064-213
64	138	8.2	360	4	US-09-906-618-213
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73	135	8.0	249	4	US-09-336-536-42
74	135	8.0	394	4	US-09-336-536-39
75	134.5	8.0	193	4	US-09-397-243D-4
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78	134.5	8.0	450	4	US-09-905-125A-320
79	134.5	8.0	450	4	US-09-906-700-320
80	134.5	8.0	450	4	US-09-906-700-320
81	134.5	8.0	450	4	US-09-903-603A-320
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86	133.5	7.9	398	4	US-09-778-510-6
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88	133.5	7.9	398	4	US-09-905-125A-84
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95	133.5	7.9	398	4	US-09-906-618-84
96	133.5	7.9	432	4	US-09-778-510-2
97	133	7.9	561	4	US-09-866-510-24
98	133	7.9	1059	4	US-09-907-794A-290
99	133	7.9	1059	4	US-09-905-125A-290
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101	133	7.9	1059	4	US-09-906-700-290	Sequence 290, App	174	120.5	7.1	529	4	US-09-823-038A-31	Sequence 31, Appl
102	133	7.9	1059	4	US-09-903-603A-290	Sequence 290, App	175	120	7.1	312	4	US-09-254-455A-9	Sequence 9, Appl
103	133	7.9	1059	4	US-09-904-920A-290	Sequence 290, App	176	120	7.1	312	4	US-09-907-794A-64	Sequence 64, Appl
104	133	7.9	1059	4	US-09-909-064-290	Sequence 290, App	177	120	7.1	312	4	US-09-905-125A-64	Sequence 64, Appl
105	133	7.9	1059	4	US-09-905-381A-290	Sequence 290, App	178	120	7.1	312	4	US-09-902-775A-64	Sequence 64, Appl
106	133	7.9	1059	4	US-09-906-610-290	Sequence 290, App	179	120	7.1	312	4	US-09-906-700-64	Sequence 64, Appl
107	133	7.9	1090	4	US-09-866-510-14	Sequence 14, Appl	180	120	7.1	312	4	US-09-903-603A-64	Sequence 64, Appl
108	133	7.9	1106	1	US-08-180-195-2	Sequence 2, Appl	181	120	7.1	312	4	US-09-904-920A-64	Sequence 64, Appl
109	133	7.9	1106	1	US-08-168-917-2	Sequence 2, Appl	182	120	7.1	312	4	US-09-909-064-64	Sequence 64, Appl
110	133	7.9	1106	1	US-08-477-329-2	Sequence 2, Appl	183	120	7.1	312	4	US-09-905-381A-64	Sequence 64, Appl
111	133	7.9	1106	2	US-08-475-458-2	Sequence 2, Appl	184	120	7.1	312	4	US-09-906-618A-64	Sequence 64, Appl
112	133	7.9	1106	2	US-08-460-510-2	Sequence 2, Appl	185	120	7.1	312	4	US-09-953-499-9	Sequence 9, Appl
113	133	7.9	1106	2	US-08-460-490-2	Sequence 2, Appl	186	120	7.1	1101	3	US-08-986-485-2	Sequence 2, Appl
114	133	7.9	1106	3	US-08-980-400-2	Sequence 2, Appl	187	119.5	7.1	298	4	US-09-152-050-76	Sequence 76, Appl
115	133	7.9	1106	3	US-08-462-728-4	Sequence 4, Appl	188	119.5	7.1	464	2	US-08-602-725-32	Sequence 32, Appl
116	133	7.9	1106	3	US-09-583-459A-2	Sequence 2, Appl	189	119.5	7.1	464	4	US-09-949-016-6116	Sequence 6116, Ap
117	133	7.9	1106	3	US-09-583-210-2	Sequence 2, Appl	190	119.5	7.1	464	4	US-09-949-016-7525	Sequence 7525, Ap
118	133	7.9	1106	3	US-09-583-449A-2	Sequence 2, Appl	191	119.5	7.1	467	3	US-09-046-736-2	Sequence 2, Appl
119	133	7.9	1106	3	US-09-435-059-2	Sequence 2, Appl	192	119	7.0	365	4	US-09-336-536-40	Sequence 40, Appl
120	133	7.9	1106	3	US-08-461-917-4	Sequence 4, Appl	193	119	7.0	1091	3	US-08-986-485-5	Sequence 5, Appl
121	133	7.9	1106	4	US-08-464-436-4	Sequence 4, Appl	194	118.5	7.0	321	6	5169835-17	Patent No. 5169835
122	133	7.9	1106	4	US-08-464-436-4	Sequence 4, Appl	195	118.5	7.0	321	6	5169835-17	Patent No. 5169835
123	133	7.9	1106	4	US-09-866-510-16	Sequence 16, Appl	196	118	7.0	518	3	US-09-240-915-8	Sequence 8, Appl
124	133	7.9	1106	4	US-09-866-510-18	Sequence 18, Appl	197	118	7.0	518	3	US-09-591-435-8	Sequence 8, Appl
125	133	7.9	1106	4	US-09-866-510-20	Sequence 20, Appl	198	118	7.0	547	1	US-08-473-981A-6	Sequence 6, Appl
126	133	7.9	1106	4	US-09-866-510-22	Sequence 22, Appl	199	118	7.0	547	2	US-08-474-087-6	Sequence 6, Appl
127	133	7.9	1106	5	PCT-US92-00730-2	Sequence 2, Appl	200	118	7.0	758	2	US-08-874-678-1	Sequence 1, Appl
128	133	7.9	1106	5	PCT-US92-00862-2	Sequence 2, Appl	201	118	7.0	758	3	US-08-643-839-1	Sequence 1, Appl
129	133	7.9	1119	4	US-09-907-794A-294	Sequence 294, App	202	118	7.0	758	3	US-09-051-363-24	Sequence 24, Appl
130	133	7.9	1119	4	US-09-905-125A-294	Sequence 294, App	203	118	7.0	758	3	US-09-348-886-1	Sequence 1, Appl
131	133	7.9	1119	4	US-09-902-775A-294	Sequence 294, App	204	118	7.0	780	1	US-08-232-538-14	Sequence 14, Appl
132	133	7.9	1119	4	US-09-906-700-294	Sequence 294, App	205	118	7.0	780	2	US-08-786-164-14	Sequence 14, Appl
133	133	7.9	1119	4	US-09-903-603A-294	Sequence 294, App	206	118	7.0	1338	3	US-08-750-141A-3	Sequence 3, Appl
134	133	7.9	1119	4	US-09-904-920A-294	Sequence 294, App	207	118	7.0	1338	4	US-09-119-014D-6	Sequence 6, Appl
135	133	7.9	1119	4	US-09-909-064-294	Sequence 294, App	208	117.5	7.0	308	2	US-08-414-657D-46	Sequence 46, Appl
136	133	7.9	1119	4	US-09-905-381A-294	Sequence 294, App	209	117.5	7.0	325	2	US-08-414-657D-2	Sequence 2, Appl
137	133	7.9	1119	4	US-09-906-618-294	Sequence 294, App	210	117.5	7.0	325	2	US-08-414-657D-41	Sequence 41, Appl
138	132.5	7.8	630	2	US-08-752-307B-14	Sequence 14, Appl	211	117.5	7.0	325	4	US-09-135-080-2	Sequence 2, Appl
139	132.5	7.8	630	3	US-09-707-802-14	Sequence 14, Appl	212	117.5	7.0	338	2	US-08-414-657D-60	Sequence 60, Appl
140	132.5	7.8	630	3	US-09-991-326-14	Sequence 14, Appl	213	117.5	7.0	338	4	US-09-135-080-8	Sequence 8, Appl
141	131	7.8	4391	4	US-10-006-011A-2	Sequence 2, Appl	214	117.5	7.0	338	4	US-09-976-594-404	Sequence 404, App
142	129	7.6	1248	4	US-09-949-016-10595	Sequence 10595, A	215	117.5	7.0	365	4	US-09-949-016-7591	Sequence 7591, Ap
143	129	7.6	1248	4	US-09-949-016-10596	Sequence 10596, A	216	117.5	7.0	615	2	US-08-752-307B-9	Sequence 9, Appl
144	127	7.5	227	4	US-09-205-258-947	Sequence 947, App	217	117.5	7.0	615	3	US-09-707-802-9	Sequence 9, Appl
145	127	7.5	462	2	US-08-752-307B-7	Sequence 7, Appl	218	117.5	7.0	615	3	US-09-991-326-9	Sequence 9, Appl
146	127	7.5	462	3	US-09-707-802-7	Sequence 7, Appl	219	117	6.9	292	4	US-09-800-729-175	Sequence 175, App
147	127	7.5	462	3	US-09-991-326-7	Sequence 7, Appl	220	117	6.9	1447	5	US-09-041-886-25	Sequence 25, Appl
148	127	7.5	465	2	US-08-752-307B-5	Sequence 5, Appl	221	117	6.9	1447	5	PCT-US94-05277-2	Sequence 2, Appl
149	127	7.5	465	3	US-09-707-802-5	Sequence 5, Appl	222	117	6.9	1745	4	US-09-800-729-89	Sequence 89, Appl
150	127	7.5	465	3	US-09-991-326-5	Sequence 5, Appl	223	116.5	6.9	315	2	US-08-414-657D-47	Sequence 47, Appl
151	126.5	7.5	924	1	US-08-481-130-28	Sequence 28, Appl	224	116.5	6.9	338	2	US-08-414-657D-42	Sequence 42, Appl
152	126.5	7.5	924	1	US-08-656-984A-28	Sequence 28, Appl	225	116.5	6.9	338	2	US-08-414-657D-43	Sequence 43, Appl
153	126.5	7.5	924	1	US-08-485-604-28	Sequence 28, Appl	226	116.5	6.9	338	4	US-09-135-080-4	Sequence 4, Appl
154	126.5	7.5	924	2	US-08-487-595-28	Sequence 28, Appl	227	116.5	6.9	607	2	US-08-752-307B-12	Sequence 12, Appl
155	124.5	7.4	387	3	US-09-175-928-2	Sequence 2, Appl	228	116.5	6.9	607	3	US-09-707-802-12	Sequence 12, Appl
156	123.5	7.3	833	4	US-09-949-016-11496	Sequence 11496, A	229	116.5	6.9	607	3	US-09-991-326-12	Sequence 12, Appl
157	123.5	7.3	1180	4	US-09-949-016-6577	Sequence 6577, Ap	230	116	6.9	1709	4	US-09-949-016-10503	Sequence 10503, A
158	123	7.3	313	4	US-09-700-397-4	Sequence 4, Appl	231	115.5	6.8	303	4	US-09-651-200-23	Sequence 23, Appl
159	123	7.3	344	4	US-09-700-397-3	Sequence 3, Appl	232	115.5	6.8	303	4	US-09-441-411-15	Sequence 15, Appl
160	122.5	7.3	588	4	US-09-949-016-10547	Sequence 10547, A	233	115.5	6.8	303	4	US-09-441-411-20	Sequence 20, Appl
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162	122.5	7.3	612	3	US-09-707-802-11	Sequence 11, Appl	235	115.5	6.8	309	3	US-08-479-744A-23	Sequence 23, Appl
163	122.5	7.3	612	3	US-09-991-326-11	Sequence 11, Appl	236	115.5	6.8	309	3	US-08-280-757B-23	Sequence 23, Appl
164	122.5	7.3	1268	3	US-08-506-296B-28	Sequence 28, Appl	237	115.5	6.8	309	3	US-08-205-697A-21	Sequence 21, Appl
165	122	7.2	611	2	US-08-752-307B-10	Sequence 10, Appl	238	115.5	6.8	309	3	US-08-702-525-21	Sequence 21, Appl
166	122	7.2	611	3	US-09-707-802-10	Sequence 10, Appl	239	115.5	6.8	309	4	US-09-651-200-22	Sequence 22, Appl
167	122	7.2	611	3	US-09-991-326-10	Sequence 10, Appl	240	115.5	6.8	309	4	US-09-667-135-33	Sequence 33, Appl
168	120.5	7.1	373	4	US-09-823-038A-60	Sequence 60, Appl	241	115.5	6.8	309	4	US-09-425-762-23	Sequence 23, Appl
169	120.5	7.1	440	4	US-09-866-028-61	Sequence 61, Appl	242	115.5	6.8	309	4	US-09-837-867A-21	Sequence 21, Appl
170	120.5	7.1	440	4	US-09-944-457-61	Sequence 61, Appl	243	115.5	6.8	309	4	US-09-206-132-4	Sequence 4, Appl
171	120.5	7.1	442	4	US-09-778-510-20	Sequence 20, Appl	244	115.5	6.8	309	4	US-09-441-411-13	Sequence 13, Appl
172	120.5	7.1	442	4	US-09-930-803-1	Sequence 1, Appl	245	115.5	6.8	309	4	US-09-441-411-18	Sequence 18, Appl
173	120.5	7.1	529	3	US-09-383-586-31	Sequence 31, Appl	246	115.5	6.8	309	4	US-09-441-411-24	Sequence 24, Appl

247	115.5	6.8	309	4	US-09-425-516-23	Sequence 23, Appl	320	112	6.6	897	4	US-08-467-602-337	Sequence 337, App
248	115.5	6.8	309	5	PCT-US95-02576-21	Sequence 21, Appl	321	112	6.6	897	4	US-08-411-295F-263	Sequence 263, App
249	115.5	6.8	314	3	US-08-205-697A-13	Sequence 13, Appl	322	112	6.6	911	4	US-08-467-602-337	Sequence 337, App
250	115.5	6.8	314	3	US-08-702-525-13	Sequence 13, Appl	323	112	6.6	911	4	US-08-411-295F-299	Sequence 299, App
251	115.5	6.8	314	3	US-09-837-867A-13	Sequence 13, Appl	324	112	6.6	931	4	US-08-467-602-379	Sequence 379, App
252	115.5	6.8	314	4	US-09-441-411-14	Sequence 14, Appl	325	112	6.6	931	4	US-08-411-295F-305	Sequence 305, App
253	115.5	6.8	314	4	US-09-441-411-19	Sequence 19, Appl	326	112	6.6	946	5	PCT-US95-08493-13	Sequence 13, Appl
254	115.5	6.8	314	5	PCT-US95-02576-13	Sequence 13, Appl	327	111.5	6.6	212	4	US-09-949-016-10458	Sequence 10458, A
255	115.5	6.8	356	4	US-09-441-411-11	Sequence 11, Appl	328	111.5	6.6	344	2	US-08-602-725-34	Sequence 34, Appl
256	115.5	6.8	356	4	US-09-441-411-12	Sequence 12, Appl	329	111	6.6	252	2	US-08-414-657D-56	Sequence 56, Appl
257	115.5	6.8	356	4	US-09-441-411-16	Sequence 16, Appl	330	111	6.6	252	2	US-08-414-657D-57	Sequence 57, Appl
258	115.5	6.8	356	4	US-09-441-411-17	Sequence 17, Appl	331	111	6.6	795	4	US-09-949-016-7119	Sequence 7119, App
259	115.5	6.8	423	4	US-09-778-510-22	Sequence 22, Appl	332	110.5	6.5	349	4	US-09-924-103-4	Sequence 4, Appl
260	115.5	6.8	738	3	US-08-478-208-32	Sequence 32, Appl	333	110.5	6.5	390	2	US-08-979-424-1	Sequence 1, Appl
261	115.5	6.8	738	3	US-09-336-536-73	Sequence 73, Appl	334	110.5	6.5	390	4	US-09-907-794A-39	Sequence 39, Appl
262	115.5	6.8	1253	3	US-08-506-296B-18	Sequence 18, Appl	335	110.5	6.5	390	4	US-09-905-125A-39	Sequence 39, Appl
263	115	6.8	315	4	US-09-910-174B-28	Sequence 28, Appl	336	110.5	6.5	390	4	US-09-902-775A-39	Sequence 39, Appl
264	115	6.8	315	4	US-09-620-461-28	Sequence 28, Appl	337	110.5	6.5	390	4	US-09-905-700-39	Sequence 39, Appl
265	115	6.8	819	4	US-09-949-016-11044	Sequence 11044, A	338	110.5	6.5	390	4	US-09-903-603A-39	Sequence 39, Appl
266	115	6.8	1209	4	US-09-130-158A-2	Sequence 2, Appl	339	110.5	6.5	390	4	US-09-904-920A-39	Sequence 39, Appl
267	114.5	6.8	431	3	US-09-038-832-2	Sequence 2, Appl	340	110.5	6.5	390	4	US-09-905-064-39	Sequence 39, Appl
268	114.5	6.8	431	3	US-09-038-832-4	Sequence 4, Appl	341	110.5	6.5	390	4	US-09-905-381A-39	Sequence 39, Appl
269	114.5	6.8	447	4	US-09-949-016-8211	Sequence 8211, App	342	110.5	6.5	390	4	US-09-905-618-39	Sequence 39, Appl
270	114.5	6.8	479	4	US-09-723-368-2	Sequence 2, Appl	343	110.5	6.5	513	4	US-09-910-174B-18	Sequence 18, Appl
271	114.5	6.8	479	4	US-09-949-016-6278	Sequence 6278, App	344	110.5	6.5	513	4	US-09-620-461-18	Sequence 18, Appl
272	114.5	6.8	504	4	US-09-949-016-7020	Sequence 7020, App	345	110.5	6.5	1070	4	US-09-961-403-3	Sequence 3, Appl
273	114.5	6.8	511	4	US-09-949-016-10054	Sequence 10054, A	346	110.5	6.5	316	4	US-09-910-174B-24	Sequence 24, Appl
274	114.5	6.8	522	4	US-09-949-016-7563	Sequence 7563, App	347	110	6.5	316	4	US-09-620-461-24	Sequence 24, Appl
275	113.5	6.7	624	4	US-08-467-602-326	Sequence 326, App	348	110	6.5	316	4	US-08-314-615-1	Sequence 1, Appl
276	113.5	6.7	624	4	US-08-411-295F-252	Sequence 252, App	349	110	6.5	547	1	US-08-314-615-1	Sequence 1, Appl
277	113.5	6.7	658	4	US-08-467-602-368	Sequence 368, App	350	110	6.5	547	1	US-08-433-010-1	Sequence 1, Appl
278	113.5	6.7	658	4	US-08-411-295F-294	Sequence 294, App	351	110	6.5	547	1	US-08-433-010-1	Sequence 1, Appl
279	113.5	6.7	841	4	US-08-467-602-327	Sequence 327, App	352	110	6.5	547	1	US-08-482-882-1	Sequence 1, Appl
280	113.5	6.7	841	4	US-08-411-295F-253	Sequence 253, App	353	110	6.5	547	2	US-08-483-389-1	Sequence 1, Appl
281	113.5	6.7	875	4	US-08-467-602-329	Sequence 329, App	354	110	6.5	547	2	US-08-487-113D-1	Sequence 1, Appl
282	113.5	6.7	875	4	US-08-411-295F-295	Sequence 295, App	355	110	6.5	547	2	US-08-473-503-1	Sequence 1, Appl
283	113.5	6.7	888	4	US-08-467-602-325	Sequence 325, App	356	110	6.5	547	2	US-08-483-932-1	Sequence 1, Appl
284	113.5	6.7	988	4	US-08-411-295F-251	Sequence 251, App	357	110	6.5	547	3	US-08-720-420A-1	Sequence 1, Appl
285	113.5	6.7	922	4	US-08-467-602-367	Sequence 367, App	358	110	6.5	547	3	US-08-714-017-1	Sequence 1, Appl
286	113.5	6.7	922	4	US-08-411-295F-293	Sequence 293, App	359	110	6.5	547	3	US-08-863-790-1	Sequence 1, Appl
287	113	6.7	922	2	US-08-414-657D-48	Sequence 48, Appl	360	110	6.5	547	3	US-08-475-680-1	Sequence 1, Appl
288	113	6.7	294	6	5260223-1	Patent No. 5260223	361	110	6.5	547	3	US-08-296-749-1	Sequence 1, Appl
289	113	6.7	294	6	5260223-1	Patent No. 5260223	362	110	6.5	547	4	US-08-314-369-1	Sequence 1, Appl
290	113	6.7	304	2	US-08-414-657D-44	Sequence 44, Appl	363	110	6.5	837	4	US-09-949-016-6515	Sequence 6515, App
291	113	6.7	497	4	US-09-499-846-6	Sequence 6, Appl	364	110	6.5	917	1	US-08-245-295-2	Sequence 2, Appl
292	113	6.7	497	4	US-09-499-846-10	Sequence 10, Appl	365	110	6.5	917	1	US-08-481-130-2	Sequence 2, Appl
293	112.5	6.7	514	4	US-09-949-016-11380	Sequence 11380, A	366	110	6.5	917	1	US-08-656-984A-2	Sequence 2, Appl
294	112.5	6.7	517	4	US-09-723-368-4	Sequence 4, Appl	367	110	6.5	917	2	US-08-483-604-2	Sequence 2, Appl
295	112	6.6	287	2	US-08-414-657D-49	Sequence 49, Appl	368	110	6.5	917	2	US-08-487-595-2	Sequence 2, Appl
296	112	6.6	310	2	US-08-414-657D-45	Sequence 45, Appl	369	109.5	6.5	488	4	US-09-499-846-12	Sequence 12, Appl
297	112	6.6	322	3	US-09-383-586-33	Sequence 33, Appl	370	109.5	6.5	526	4	US-09-910-174B-9	Sequence 9, Appl
298	112	6.6	322	4	US-09-823-038A-33	Sequence 33, Appl	371	109.5	6.5	526	4	US-09-620-461-9	Sequence 9, Appl
299	112	6.6	613	3	US-08-470-335-230	Sequence 230, App	372	109.5	6.5	526	4	US-09-949-016-6122	Sequence 6122, App
300	112	6.6	613	4	US-08-467-602-329	Sequence 329, App	373	109.5	6.5	540	4	US-09-949-016-11644	Sequence 11644, A
301	112	6.6	613	4	US-08-411-295F-255	Sequence 255, App	374	109	6.5	339	4	US-09-719-243-2	Sequence 2, Appl
302	112	6.6	633	4	US-08-467-602-335	Sequence 335, App	375	109	6.5	358	4	US-09-719-243-3	Sequence 3, Appl
303	112	6.6	633	4	US-08-411-295F-261	Sequence 261, App	376	109	6.5	486	2	US-08-432-016-6	Sequence 6, Appl
304	112	6.6	647	4	US-08-467-602-371	Sequence 371, App	377	109	6.5	486	2	US-08-684-594-6	Sequence 6, Appl
305	112	6.6	647	4	US-08-411-295F-297	Sequence 297, App	378	109	6.5	525	4	US-09-499-846-4	Sequence 4, Appl
306	112	6.6	667	4	US-08-467-602-377	Sequence 377, App	379	109	6.5	525	4	US-09-499-846-8	Sequence 8, Appl
307	112	6.6	667	4	US-08-411-295F-303	Sequence 303, App	380	109	6.5	622	4	US-09-499-846-2	Sequence 2, Appl
308	112	6.6	830	3	US-08-470-335-231	Sequence 331, App	381	109	6.5	623	4	US-09-949-016-11206	Sequence 11206, A
309	112	6.6	830	4	US-08-467-602-330	Sequence 330, App	382	109	6.5	646	4	US-09-949-016-6728	Sequence 6728, App
310	112	6.6	830	4	US-08-411-295F-256	Sequence 256, App	383	109	6.5	646	4	US-09-653-961-2	Sequence 2, Appl
311	112	6.6	850	4	US-08-467-602-336	Sequence 336, App	384	109	6.5	646	4	US-09-653-961-4	Sequence 4, Appl
312	112	6.6	850	4	US-08-411-295F-262	Sequence 262, App	385	109	6.5	668	4	US-09-949-016-8139	Sequence 8139, App
313	112	6.6	864	4	US-08-467-602-372	Sequence 372, App	386	108.5	6.4	738	6	5264554-2	Patent No. 5264554
314	112	6.6	864	4	US-08-411-295F-298	Sequence 298, App	387	108.5	6.4	738	6	5264554-2	Patent No. 5264554
315	112	6.6	877	3	US-08-470-335-232	Sequence 332, App	388	108	6.4	421	3	US-08-659-984A-1	Sequence 1, Appl
316	112	6.6	877	4	US-08-467-602-331	Sequence 331, App	389	108	6.4	421	3	US-08-660-531-1	Sequence 1, Appl
317	112	6.6	877	4	US-08-411-295F-257	Sequence 257, App	390	108	6.4	444	2	US-08-659-984A-5	Sequence 5, Appl
318	112	6.6	884	4	US-08-467-602-378	Sequence 378, App	391	108	6.4	444	3	US-08-660-531-5	Sequence 5, Appl
319	112	6.6	884	4	US-08-411-295F-304	Sequence 304, App	392	108	6.4	868	1	US-08-374-834-1	Sequence 1, Appl

393	108	6.4	868	2	US-08-644-271-1	Sequence 1, Appli	466	105	6.2	1509	4	US-09-677-046A-2	Sequence 2, Appli
394	108	6.4	868	4	US-09-077-955-1	Sequence 1, Appli	467	104.5	6.2	1461	4	US-09-976-594-531	Sequence 531, App
395	107.5	6.4	290	4	US-09-910-174B-19	Sequence 19, Appl	468	104	6.2	307	4	US-09-197-970B-3	Sequence 3, Appli
396	107.5	6.4	290	4	US-09-620-461-19	Sequence 19, Appl	469	104	6.2	307	4	US-08-467-602-221	Sequence 221, App
397	107.5	6.4	340	4	US-09-651-200-2	Sequence 2, Appli	470	104	6.2	409	4	US-08-411-295F-147	Sequence 147, App
398	107.5	6.4	350	4	US-09-651-200-25	Sequence 25, Appl	471	104	6.2	443	4	US-08-467-602-263	Sequence 263, App
399	107.5	6.4	350	4	US-09-10-174B-17	Sequence 17, Appl	472	104	6.2	443	4	US-08-411-295F-189	Sequence 189, App
400	107.5	6.4	350	4	US-09-620-461-17	Sequence 17, Appl	473	104	6.2	478	5	PCT-US95-08493-15	Sequence 15, Appl
401	107.5	6.4	441	4	US-09-651-200-4	Sequence 4, Appli	474	104	6.2	626	4	US-08-467-602-222	Sequence 222, App
402	107.5	6.4	1297	3	US-09-540-245A-17	Sequence 17, Appl	475	104	6.2	626	4	US-08-411-295F-148	Sequence 148, App
403	107.5	6.4	1311	1	US-08-340-011-5	Sequence 5, Appli	476	104	6.2	658	4	US-08-467-602-305	Sequence 305, App
404	107.5	6.4	1311	3	US-08-901-710-5	Sequence 5, Appli	477	104	6.2	658	4	US-08-411-295F-231	Sequence 231, App
405	107.5	6.4	1311	4	US-09-169-079-5	Sequence 5, Appli	478	104	6.2	660	4	US-08-467-602-264	Sequence 264, App
406	107	6.3	624	2	US-08-642-406A-22	Sequence 22, Appl	479	104	6.2	660	4	US-08-411-295F-190	Sequence 190, App
407	107	6.3	624	3	US-09-199-534-22	Sequence 22, Appl	480	104	6.2	673	4	US-08-467-602-220	Sequence 220, App
408	107	6.3	624	4	US-09-199-534-22	Sequence 22, Appl	481	104	6.2	673	4	US-08-411-295F-146	Sequence 146, App
409	107	6.3	773	3	US-08-434-000A-2	Sequence 2, Appli	482	104	6.2	692	4	US-08-467-602-347	Sequence 347, App
410	107	6.3	773	3	US-09-312-157-2	Sequence 2, Appli	483	104	6.2	692	4	US-08-411-295F-273	Sequence 273, App
411	107	6.3	773	4	US-09-717-888-2	Sequence 2, Appli	484	104	6.2	707	4	US-08-467-602-262	Sequence 262, App
412	106.5	6.3	731	1	US-07-921-807B-5	Sequence 5, Appli	485	104	6.2	707	4	US-08-411-295F-188	Sequence 188, App
413	106.5	6.3	731	1	US-08-441-944A-5	Sequence 5, Appli	486	104	6.2	860	5	PCT-US95-08493-19	Sequence 19, Appl
414	106.5	6.3	731	3	US-08-439-992A-3	Sequence 3, Appli	487	104	6.2	868	5	PCT-US95-08493-21	Sequence 21, Appl
415	106	6.3	729	1	US-07-640-029-3	Sequence 3, Appli	488	104	6.2	875	4	US-08-467-602-306	Sequence 306, App
416	106	6.3	1474	4	US-09-677-046A-4	Sequence 4, Appli	489	104	6.2	875	4	US-08-411-295F-232	Sequence 232, App
417	106	6.3	1953	4	US-09-917-254-92	Sequence 92, Appl	490	104	6.2	909	4	US-08-467-602-348	Sequence 348, App
418	105.5	6.2	246	1	US-07-843-125-11	Sequence 11, Appl	491	104	6.2	909	4	US-08-411-295F-274	Sequence 274, App
419	105.5	6.2	351	5	PCT-US93-05703-2	Sequence 2, Appli	492	104	6.2	922	4	US-08-467-602-304	Sequence 304, App
420	105.5	6.2	408	3	US-09-724-864-62	Sequence 62, Appli	493	104	6.2	922	4	US-08-411-295F-230	Sequence 230, App
421	105.5	6.2	589	2	US-08-724-394A-1	Sequence 1, Appli	494	104	6.2	956	4	US-08-467-602-346	Sequence 346, App
422	105.5	6.2	604	3	US-08-470-335-227	Sequence 227, App	495	104	6.2	956	4	US-08-411-295F-272	Sequence 272, App
423	105.5	6.2	604	4	US-08-467-602-318	Sequence 318, App	496	104	6.2	1503	4	US-09-677-046A-6	Sequence 6, Appli
424	105.5	6.2	604	4	US-08-411-295F-244	Sequence 244, App	497	103.5	6.1	534	4	US-09-651-200-6	Sequence 6, Appli
425	105.5	6.2	609	4	US-09-949-016-7747	Sequence 7747, Ap	498	103.5	6.1	534	4	US-09-651-200-24	Sequence 24, Appl
426	105.5	6.2	609	4	US-09-949-016-7748	Sequence 7748, Ap	499	103.5	6.1	640	4	US-09-949-016-7565	Sequence 7565, Ap
427	105.5	6.2	609	4	US-09-949-016-7749	Sequence 7749, Ap	500	103.5	6.1	668	1	US-08-232-538-13	Sequence 13, Appl
428	105.5	6.2	609	4	US-09-949-016-7750	Sequence 7750, Ap	501	103.5	6.1	668	2	US-08-786-164-13	Sequence 13, Appl
429	105.5	6.2	609	4	US-09-949-016-7751	Sequence 7751, Ap	502	103.5	6.1	764	3	US-09-142-956B-14	Sequence 14, Appl
430	105.5	6.2	609	4	US-09-949-016-7752	Sequence 7752, Ap	503	103.5	6.1	767	2	US-08-874-678-2	Sequence 2, Appli
431	105.5	6.2	609	4	US-09-949-016-7753	Sequence 7753, Ap	504	103.5	6.1	767	3	US-08-643-839-2	Sequence 2, Appli
432	105.5	6.2	609	4	US-09-949-016-7754	Sequence 7754, Ap	505	103.5	6.1	767	3	US-03-348-886-2	Sequence 2, Appli
433	105.5	6.2	638	4	US-08-467-602-360	Sequence 360, App	506	103.5	6.1	788	1	US-08-232-538-15	Sequence 15, Appl
434	105.5	6.2	638	4	US-08-411-295F-286	Sequence 286, App	507	103.5	6.1	788	2	US-08-786-164-15	Sequence 15, Appl
435	105.5	6.2	651	4	US-09-270-767-44877	Sequence 44877, A	508	103.5	6.1	1356	1	US-08-810-116-8	Sequence 8, Appli
436	105.5	6.2	733	1	US-07-640-029-4	Sequence 4, Appli	509	103.5	6.1	1356	2	US-07-930-548A-8	Sequence 8, Appli
437	105.5	6.2	733	1	US-07-921-807B-6	Sequence 6, Appli	510	103.5	6.1	1356	3	US-09-098-707A-2	Sequence 2, Appli
438	105.5	6.2	733	1	US-08-441-944A-6	Sequence 6, Appli	511	103.5	6.1	1356	3	US-03-483-539-2	Sequence 2, Appli
439	105.5	6.2	733	3	US-08-439-992A-4	Sequence 4, Appli	512	103.5	6.1	1356	3	US-09-949-016-6198	Sequence 6198, Ap
440	105.5	6.2	817	1	US-07-640-029-2	Sequence 2, Appli	513	103.5	6.1	1381	3	US-09-540-245A-16	Sequence 16, Appl
441	105.5	6.2	820	1	US-07-921-807B-3	Sequence 3, Appli	514	103.5	6.1	1456	4	US-09-949-016-9853	Sequence 9853, Ap
442	105.5	6.2	820	3	US-08-441-944A-3	Sequence 3, Appli	515	103.5	6.1	1651	3	US-09-540-245A-18	Sequence 18, Appl
443	105.5	6.2	820	1	US-08-439-992A-1	Sequence 1, Appli	516	103	6.1	240	2	US-07-956-399-2	Sequence 2, Appli
444	105.5	6.2	821	3	US-08-470-335-228	Sequence 228, App	517	103	6.1	806	3	US-03-383-630-3	Sequence 3, Appli
445	105.5	6.2	821	4	US-08-467-602-319	Sequence 319, App	518	102.5	6.1	326	1	US-08-225-477B-6	Sequence 6, Appli
446	105.5	6.2	821	4	US-08-411-295F-245	Sequence 245, App	519	102.5	6.1	326	5	PCT-US95-04353-6	Sequence 6, Appli
447	105.5	6.2	822	1	US-07-997-133-1	Sequence 1, Appli	520	102.5	6.1	398	4	US-08-467-602-224	Sequence 224, App
448	105.5	6.2	822	1	US-07-921-807B-4	Sequence 4, Appli	521	102.5	6.1	398	4	US-08-411-295F-150	Sequence 150, App
449	105.5	6.2	822	1	US-08-459-296-2	Sequence 2, Appli	522	102.5	6.1	418	4	US-08-467-602-230	Sequence 230, App
450	105.5	6.2	822	1	US-08-441-944A-4	Sequence 4, Appli	523	102.5	6.1	418	4	US-08-411-295F-156	Sequence 156, App
451	105.5	6.2	822	1	US-08-451-822A-12	Sequence 12, Appl	524	102.5	6.1	432	4	US-08-467-602-266	Sequence 266, App
452	105.5	6.2	822	3	US-08-439-992A-2	Sequence 2, Appli	525	102.5	6.1	432	4	US-08-411-295F-192	Sequence 192, App
453	105.5	6.2	822	3	US-08-323-430-12	Sequence 12, Appl	526	102.5	6.1	452	4	US-08-467-602-272	Sequence 272, App
454	105.5	6.2	855	4	US-08-467-602-361	Sequence 361, App	527	102.5	6.1	452	4	US-08-411-295F-198	Sequence 198, App
455	105.5	6.2	855	4	US-08-411-295F-287	Sequence 287, App	528	102.5	6.1	458	4	US-09-435-956A-1	Sequence 1, Appli
456	105.5	6.2	868	3	US-08-470-335-229	Sequence 229, App	529	102.5	6.1	615	4	US-08-467-602-225	Sequence 225, App
457	105.5	6.2	868	4	US-08-467-602-317	Sequence 317, App	530	102.5	6.1	615	4	US-08-411-295F-151	Sequence 151, App
458	105.5	6.2	868	4	US-08-411-295F-243	Sequence 243, App	531	102.5	6.1	635	4	US-08-467-602-231	Sequence 231, App
459	105.5	6.2	869	1	US-08-374-834-16	Sequence 16, Appl	532	102.5	6.1	635	4	US-08-411-295F-157	Sequence 157, App
460	105.5	6.2	869	2	US-08-644-271-29	Sequence 29, Appl	533	102.5	6.1	647	3	US-08-470-335-243	Sequence 243, App
461	105.5	6.2	869	4	US-09-077-955-33	Sequence 33, Appl	534	102.5	6.1	647	4	US-08-467-602-308	Sequence 308, App
462	105.5	6.2	902	4	US-08-467-602-359	Sequence 359, App	535	102.5	6.1	647	4	US-08-411-295F-234	Sequence 234, App
463	105.5	6.2	902	4	US-08-411-295F-285	Sequence 285, App	536	102.5	6.1	649	4	US-08-467-602-267	Sequence 267, App
464	105	6.2	540	2	US-08-724-394A-4	Sequence 4, Appli	537	102.5	6.1	649	4	US-08-411-295F-193	Sequence 193, App
465	105	6.2	828	1	US-08-261-304-2	Sequence 2, Appli	538	102.5	6.1	662	4	US-08-467-602-226	Sequence 226, App

539	102.5	6.1	662	4	US-08-411-295F-152	Sequence 152, App	612	100.5	6.0	626	4	US-08-411-295F-211	Sequence 211, App
540	102.5	6.1	667	4	US-08-467-602-314	Sequence 314, App	613	100.5	6.0	638	3	US-08-470-335-240	Sequence 240, App
541	102.5	6.1	667	4	US-08-411-295F-240	Sequence 240, App	614	100.5	6.0	638	4	US-08-467-602-237	Sequence 297, App
542	102.5	6.1	669	4	US-08-467-602-273	Sequence 273, App	615	100.5	6.0	638	4	US-08-411-295F-223	Sequence 223, App
543	102.5	6.1	669	4	US-08-411-295F-199	Sequence 199, App	616	100.5	6.0	672	4	US-08-467-602-339	Sequence 339, App
544	102.5	6.1	681	4	US-08-467-602-350	Sequence 350, App	617	100.5	6.0	672	4	US-08-411-295F-265	Sequence 265, App
545	102.5	6.1	681	4	US-08-411-295F-276	Sequence 276, App	618	100.5	6.0	673	4	US-08-467-602-283	Sequence 283, App
546	102.5	6.1	682	4	US-08-467-602-232	Sequence 232, App	619	100.5	6.0	673	4	US-08-411-295F-209	Sequence 209, App
547	102.5	6.1	682	4	US-08-411-295F-158	Sequence 158, App	620	100.5	6.0	802	3	US-09-173-151A-33	Sequence 33, Appl
548	102.5	6.1	696	4	US-08-467-602-268	Sequence 268, App	621	100.5	6.0	855	3	US-08-470-335-241	Sequence 241, App
549	102.5	6.1	696	4	US-08-411-295F-194	Sequence 194, App	622	100.5	6.0	855	4	US-08-467-602-298	Sequence 298, App
550	102.5	6.1	701	4	US-08-467-602-356	Sequence 356, App	623	100.5	6.0	855	4	US-08-411-295F-224	Sequence 224, App
551	102.5	6.1	701	4	US-08-411-295F-282	Sequence 282, App	624	100.5	6.0	869	4	US-09-715-249-8	Sequence 8, Appl
552	102.5	6.1	716	4	US-08-467-602-274	Sequence 274, App	625	100.5	6.0	889	4	US-08-467-602-340	Sequence 340, App
553	102.5	6.1	716	4	US-08-411-295F-200	Sequence 200, App	626	100.5	6.0	889	4	US-08-411-295F-266	Sequence 266, App
554	102.5	6.1	864	3	US-08-470-335-244	Sequence 244, App	627	100.5	6.0	902	3	US-08-470-335-242	Sequence 242, App
555	102.5	6.1	864	4	US-08-467-602-309	Sequence 309, App	628	100.5	6.0	902	4	US-08-467-602-296	Sequence 296, App
556	102.5	6.1	864	4	US-08-411-295F-235	Sequence 235, App	629	100.5	6.0	902	4	US-08-411-295F-222	Sequence 222, App
557	102.5	6.1	884	4	US-08-467-602-315	Sequence 315, App	630	100.5	6.0	936	4	US-08-467-602-338	Sequence 338, App
558	102.5	6.1	884	4	US-08-411-295F-241	Sequence 241, App	631	100.5	6.0	936	4	US-08-411-295F-264	Sequence 264, App
559	102.5	6.1	898	4	US-08-467-602-351	Sequence 351, App	632	100	5.9	63	4	US-09-397-243D-10	Sequence 10, Appl
560	102.5	6.1	898	4	US-08-411-295F-277	Sequence 277, App	633	100	5.9	63	4	US-09-397-243D-10	Sequence 10, Appl
561	102.5	6.1	911	3	US-08-470-335-245	Sequence 245, App	634	100	5.9	124	4	US-09-858-664A-28	Sequence 28, Appl
562	102.5	6.1	911	4	US-08-467-602-310	Sequence 310, App	635	100	5.9	124	4	US-10-274-978-29	Sequence 29, Appl
563	102.5	6.1	911	4	US-08-411-295F-236	Sequence 236, App	636	100	5.9	124	4	US-10-697-263-29	Sequence 29, Appl
564	102.5	6.1	918	4	US-08-467-602-357	Sequence 357, App	637	100	5.9	322	4	US-09-667-135-2	Sequence 2, Appl
565	102.5	6.1	918	4	US-08-411-295F-283	Sequence 283, App	638	100	5.9	322	4	US-09-910-174B-29	Sequence 29, Appl
566	102.5	6.1	931	4	US-08-467-602-316	Sequence 316, App	639	100	5.9	322	4	US-09-620-461-29	Sequence 29, Appl
567	102.5	6.1	931	4	US-08-411-295F-242	Sequence 242, App	640	100	5.9	373	4	US-09-943-016-10485	Sequence 10485, A
568	102.5	6.1	945	4	US-08-467-602-352	Sequence 352, App	641	100	5.9	458	4	US-10-092-138A-25	Sequence 25, Appl
569	102.5	6.1	945	4	US-08-411-295F-278	Sequence 278, App	642	99.5	5.9	244	4	US-09-244-369B-1	Sequence 1, Appl
570	102.5	6.1	965	4	US-08-467-602-358	Sequence 358, App	643	99.5	5.9	244	4	US-09-940-391-1	Sequence 1, Appl
571	102.5	6.1	965	4	US-08-411-295F-284	Sequence 284, App	644	99.5	5.9	596	2	US-08-752-307B-13	Sequence 13, Appl
572	102.5	6.1	1184	2	US-08-918-914-1	Sequence 1, Appl	645	99.5	5.9	596	3	US-09-707-802-13	Sequence 13, Appl
573	102.5	6.1	1184	3	US-08-996-083-3	Sequence 3, Appl	646	98.5	5.9	596	3	US-09-991-326-13	Sequence 13, Appl
574	102	6.0	319	4	US-09-910-174B-12	Sequence 12, Appl	647	99.5	5.9	610	3	US-08-470-335-236	Sequence 236, App
575	102	6.0	319	4	US-09-620-461-12	Sequence 12, Appl	648	99.5	5.9	610	4	US-08-467-602-332	Sequence 332, App
576	102	6.0	347	4	US-09-667-135-4	Sequence 4, Appl	649	99.5	5.9	626	4	US-08-411-295F-258	Sequence 258, App
577	102	6.0	357	4	US-09-910-174B-14	Sequence 14, Appl	650	99.5	5.9	626	4	US-09-949-016-6213	Sequence 6213, Ap
578	102	6.0	357	4	US-09-620-461-14	Sequence 14, Appl	651	99.5	5.9	644	4	US-08-467-602-370	Sequence 374, App
579	102	6.0	642	1	US-08-217-299-1	Sequence 1, Appl	652	98.5	5.9	644	4	US-08-411-295F-300	Sequence 300, App
580	102	6.0	698	2	US-08-602-725-36	Sequence 36, Appl	653	99.5	5.9	664	4	US-09-949-016-7850	Sequence 7850, Ap
581	102	6.0	702	4	US-09-949-016-6484	Sequence 6484, Ap	654	99.5	5.9	827	3	US-08-470-335-237	Sequence 237, App
582	102	6.0	734	2	US-08-389-459A-17	Sequence 17, Appl	655	99.5	5.9	827	4	US-08-467-602-333	Sequence 333, App
583	102	6.0	734	3	US-08-987-867A-17	Sequence 17, Appl	656	99.5	5.9	827	4	US-08-411-295F-259	Sequence 259, App
584	102	6.0	740	4	US-09-949-016-8168	Sequence 8168, Ap	657	99.5	5.9	861	4	US-08-467-602-375	Sequence 375, App
585	102	6.0	1439	4	US-09-134-000C-6133	Sequence 6133, Ap	658	98.5	5.9	861	4	US-08-411-295F-301	Sequence 301, App
586	101.5	6.0	417	4	US-09-949-016-6729	Sequence 6729, Ap	659	99.5	5.9	874	3	US-08-470-335-238	Sequence 238, App
587	101.5	6.0	456	4	US-09-949-016-7564	Sequence 7564, Ap	660	99.5	5.9	874	4	US-08-467-602-334	Sequence 334, App
588	101.5	6.0	816	4	US-09-949-016-10904	Sequence 10904, A	661	99.5	5.9	874	4	US-08-411-295F-260	Sequence 260, App
589	101.5	6.0	1260	3	US-08-506-296B-21	Sequence 21, Appl	662	99.5	5.9	908	4	US-08-467-602-376	Sequence 376, App
590	101	6.0	318	6	5223394-11	Patent No. 5223394	663	99.5	5.9	908	4	US-08-411-295F-302	Sequence 302, App
591	101	6.0	318	6	5223394-11	Patent No. 5223394	664	99	5.9	336	4	US-09-949-016-7714	Sequence 7714, Ap
592	101	6.0	394	3	US-08-466-368-2	Sequence 2, Appl	665	99	5.9	336	4	US-09-949-016-7715	Sequence 7715, Ap
593	101	6.0	394	4	US-08-328-500-2	Sequence 2, Appl	666	99	5.9	341	4	US-09-336-536-28	Sequence 29, Appl
594	101	6.0	458	3	US-09-517-605-3	Sequence 3, Appl	667	99	5.9	370	4	US-09-336-536-28	Sequence 28, Appl
595	101	6.0	801	3	US-09-383-630-6	Sequence 6, Appl	668	99	5.9	374	3	US-09-046-736-4	Sequence 4, Appl
596	101	6.0	890	1	US-08-445-640-2	Sequence 2, Appl	669	99	5.9	375	4	US-08-467-602-242	Sequence 242, App
597	101	6.0	890	3	US-08-170-558-2	Sequence 2, Appl	670	99	5.9	375	4	US-08-411-295F-168	Sequence 168, App
598	101	6.0	890	3	US-08-447-314-2	Sequence 2, Appl	671	99	5.9	388	1	US-08-445-640-6	Sequence 6, Appl
599	101	6.0	890	3	US-08-445-461-2	Sequence 2, Appl	672	99	5.9	388	3	US-08-170-558-6	Sequence 6, Appl
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602	101	6.0	911	2	US-08-441-104A-1	Sequence 1, Appl	675	99	5.9	388	4	US-09-223-490-6	Sequence 6, Appl
603	101	6.0	911	2	US-08-440-816A-1	Sequence 1, Appl	676	99	5.9	394	6	5223418-2	Patent No. 5223418
604	101	6.0	911	3	US-09-417-381A-1	Sequence 1, Appl	677	99	5.9	394	6	5223418-2	Patent No. 5223418
605	100.5	6.0	304	3	US-08-862-124-14	Sequence 14, Appl	678	99	5.9	398	4	US-08-467-602-287	Sequence 287, App
606	100.5	6.0	409	4	US-08-467-602-284	Sequence 284, App	679	99	5.9	398	4	US-08-411-295F-213	Sequence 213, App
607	100.5	6.0	409	4	US-08-411-295F-210	Sequence 210, App	680	99	5.9	418	4	US-08-467-602-293	Sequence 293, App
608	100.5	6.0	455	4	US-09-949-016-6949	Sequence 6949, Ap	681	99	5.9	418	4	US-08-411-295F-219	Sequence 219, App
609	100.5	6.0	455	4	US-09-949-016-11026	Sequence 11026, A	682	99	5.9	458	3	US-09-039-555B-15	Sequence 15, Appl
610	100.5	6.0	501	2	US-08-408-095-31	Sequence 31, Appl	683	99	5.9	584	4	US-09-910-174B-16	Sequence 16, Appl
611	100.5	6.0	626	4	US-08-467-602-285	Sequence 285, App	684	99	5.9	584	4	US-09-620-461-16	Sequence 16, Appl

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686	99	5.9	592	4	US-08-411-295F-169	Sequence 169, App	759	97	5.7	532	5	PCT-US92-01785-6	Sequence 6, Appli
687	99	5.9	615	4	US-08-467-602-286	Sequence 286, App	760	97	5.7	532	5	PCT-US95-00454-6	Sequence 6, Appli
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690	99	5.9	635	4	US-08-411-295F-220	Sequence 220, App	763	97	5.7	575	3	US-09-218-950-4	Sequence 4, Appli
691	99	5.9	639	4	US-08-467-602-241	Sequence 241, App	764	97	5.7	575	4	US-08-394-388A-4	Sequence 4, Appli
692	99	5.9	639	4	US-08-411-295F-167	Sequence 167, App	765	97	5.7	575	5	PCT-US92-01785-4	Sequence 4, Appli
693	99	5.9	662	4	US-08-467-602-289	Sequence 289, App	766	97	5.7	575	5	PCT-US95-00454-4	Sequence 4, Appli
694	99	5.9	662	4	US-08-411-295F-215	Sequence 215, App	767	97	5.7	601	3	US-08-470-335-233	Sequence 233, App
695	99	5.9	682	4	US-08-467-602-295	Sequence 295, App	768	97	5.7	601	4	US-08-467-602-323	Sequence 323, App
696	99	5.9	682	4	US-08-411-295F-221	Sequence 221, App	769	97	5.7	601	4	US-08-411-295F-249	Sequence 249, App
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698	98.5	5.8	324	4	US-09-620-461-6	Sequence 6, Appli	771	97	5.7	635	4	US-08-467-602-365	Sequence 365, App
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700	98.5	5.8	434	2	US-08-720-420A-120	Sequence 120, App	773	97	5.7	662	1	US-08-261-304-7	Sequence 7, Appli
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704	98.5	5.8	602	1	US-08-339-517-5	Sequence 5, Appli	777	97	5.7	739	5	PCT-US93-00031-9	Sequence 9, Appli
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707	98	5.8	243	1	US-08-230-843-4	Sequence 4, Appli	780	97	5.7	818	3	US-08-470-335-234	Sequence 234, App
708	98	5.8	243	2	US-08-636-936-4	Sequence 4, Appli	781	97	5.7	818	4	US-08-467-602-321	Sequence 321, App
709	98	5.8	246	4	US-09-336-536-31	Sequence 31, Appli	782	97	5.7	852	4	US-08-411-295F-247	Sequence 247, App
710	98	5.8	295	6	5223394-9	Patent No. 5223394	783	97	5.7	852	4	US-08-467-602-363	Sequence 363, App
711	98	5.8	295	6	5223394-9	Patent No. 5223394	784	97	5.7	852	4	US-08-411-295F-289	Sequence 289, App
712	98	5.8	874	2	US-08-456-647B-6	Sequence 6, Appli	785	97	5.7	865	3	US-08-470-335-235	Sequence 235, App
713	98	5.8	874	2	US-08-237-401A-6	Sequence 6, Appli	786	97	5.7	865	4	US-08-467-602-322	Sequence 322, App
714	98	5.8	880	1	US-08-445-640-10	Sequence 10, Appli	787	97	5.7	865	4	US-08-411-295F-248	Sequence 248, App
715	98	5.8	880	3	US-08-170-558-10	Sequence 10, Appli	788	97	5.7	899	4	US-08-467-602-364	Sequence 364, App
716	98	5.8	880	3	US-08-447-314-10	Sequence 10, Appli	789	97	5.7	899	4	US-08-411-295F-290	Sequence 290, App
717	98	5.8	880	3	US-08-445-461-10	Sequence 10, Appli	790	97	5.7	969	4	US-09-949-016-8059	Sequence 8059, Ap
718	98	5.8	1021	1	US-09-223-490-10	Sequence 10, Appli	791	97	5.7	1367	1	US-07-813-593-4	Sequence 4, Appli
719	98	5.8	1021	1	US-08-497-025-3	Sequence 3, Appli	792	97	5.7	1367	1	US-07-977-451-6	Sequence 6, Appli
720	97.5	5.8	97	4	US-09-858-664A-27	Sequence 27, Appli	793	97	5.7	1367	1	US-07-946-507-4	Sequence 4, Appli
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722	97.5	5.8	97	4	US-10-697-263-28	Sequence 28, Appli	795	97	5.7	1367	1	US-07-906-397A-6	Sequence 6, Appli
723	97.5	5.8	246	1	US-08-197-834-7	Sequence 7, Appli	796	97	5.7	1367	1	US-08-601-831-6	Sequence 6, Appli
724	97.5	5.8	342	2	US-08-724-394A-6	Sequence 6, Appli	797	97	5.7	1367	2	US-08-443-861-2	Sequence 2, Appli
725	97.5	5.8	364	4	US-08-467-602-245	Sequence 245, App	798	97	5.7	1367	2	US-09-021-324-6	Sequence 6, Appli
726	97.5	5.8	364	4	US-08-411-295F-171	Sequence 171, App	799	97	5.7	1367	3	US-08-193-829B-2	Sequence 2, Appli
727	97.5	5.8	384	4	US-08-467-602-251	Sequence 251, App	800	97	5.7	1367	4	US-09-872-136B-6	Sequence 6, Appli
728	97.5	5.8	384	4	US-08-411-295F-177	Sequence 177, App	801	97	5.7	1367	5	PCT-US92-02750-8	Sequence 8, Appli
729	97.5	5.8	581	4	US-08-467-602-246	Sequence 246, App	802	97	5.7	1367	5	PCT-US92-05401-6	Sequence 6, Appli
730	97.5	5.8	581	4	US-08-411-295F-172	Sequence 172, App	803	97	5.7	1367	5	PCT-US92-09893-6	Sequence 6, Appli
731	97.5	5.8	601	4	US-08-467-602-252	Sequence 252, App	804	96.5	5.7	158	3	US-09-560-639-13	Sequence 13, Appli
732	97.5	5.8	601	4	US-08-467-602-252	Sequence 2, App	805	96.5	5.7	263	4	US-09-949-016-8484	Sequence 8484, Ap
733	97.5	5.8	602	1	US-08-168-091A-2	Sequence 2, Appli	806	96.5	5.7	332	4	US-09-949-016-8483	Sequence 8483, Ap
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735	97.5	5.8	628	4	US-08-411-295F-173	Sequence 173, App	808	96.5	5.7	439	4	US-09-823-038A-32	Sequence 32, Appli
736	97.5	5.8	648	4	US-08-467-602-253	Sequence 253, App	809	96.5	5.7	489	4	US-09-667-135-30	Sequence 30, Appli
737	97.5	5.8	648	4	US-08-411-295F-179	Sequence 179, App	810	96.5	5.7	560	4	US-09-949-016-8293	Sequence 21, Appli
738	97.5	5.8	821	1	US-08-339-578-2	Sequence 2, Appli	811	96.5	5.7	782	4	US-09-684-708A-21	Sequence 21, Appli
739	97	5.7	340	3	US-09-188-930-184	Sequence 184, App	812	96.5	5.7	820	1	US-08-166-717D-6	Sequence 6, Appli
740	97	5.7	340	4	US-09-312-283C-184	Sequence 29, App	813	96	5.7	389	4	US-08-467-602-213	Sequence 213, App
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742	97	5.7	398	3	US-09-218-950-29	Sequence 29, App	815	96	5.7	423	4	US-08-467-602-255	Sequence 255, App
743	97	5.7	398	4	US-08-394-388A-29	Sequence 29, Appli	816	96	5.7	423	4	US-08-411-295F-181	Sequence 181, App
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745	97	5.7	402	3	US-08-457-918-1	Sequence 1, Appli	818	96	5.7	469	3	US-09-398-496-8	Sequence 8, Appli
746	97	5.7	402	4	US-10-157-408-1	Sequence 1, Appli	819	96	5.7	606	4	US-08-467-602-214	Sequence 214, App
747	97	5.7	457	4	US-08-328-500-9	Sequence 9, Appli	820	96	5.7	606	4	US-08-411-295F-140	Sequence 140, App
748	97	5.7	458	3	US-08-466-368-4	Sequence 4, Appli	821	96	5.7	625	1	US-07-847-743B-26	Sequence 26, Appli
749	97	5.7	462	2	US-08-417-495-5	Sequence 5, Appli	822	96	5.7	625	1	US-08-456-201-26	Sequence 26, Appli
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751	97	5.7	462	3	US-09-218-950-5	Sequence 5, Appli	824	96	5.7	625	5	PCT-US92-04295A-26	Sequence 26, Appli
752	97	5.7	462	4	US-08-394-388A-5	Sequence 5, Appli	825	96	5.7	640	4	US-08-467-602-256	Sequence 256, App
753	97	5.7	462	5	PCT-US92-01785-5	Sequence 5, Appli	826	96	5.7	640	4	US-09-907-794A-292	Sequence 292, App
754	97	5.7	462	5	PCT-US95-00454-5	Sequence 5, Appli	827	96	5.7	640	4	US-09-905-125A-292	Sequence 292, App
755	97	5.7	532	2	US-08-417-495-6	Sequence 6, Appli	828	96	5.7	640	4	US-09-902-775A-292	Sequence 292, App
756	97	5.7	532	2	US-08-284-391B-6	Sequence 6, Appli	829	96	5.7	640	4	US-09-906-700-292	Sequence 292, App
757	97	5.7	532	3	US-09-218-950-6	Sequence 6, Appli	830	96	5.7	640	4	US-08-411-295F-182	Sequence 182, App

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832	96	5.7	640	4	US-09-904-920A-292	Sequence 292, App	905	94.5	5.6	999	4	US-09-949-016-6718	Sequence 6718, Ap
833	96	5.7	640	4	US-09-909-064-292	Sequence 292, App	906	94	5.6	198	4	US-09-569-611C-34	Sequence 34, Appl
834	96	5.7	640	4	US-09-905-381A-292	Sequence 292, App	907	94	5.6	203	4	US-09-270-767-60345	Sequence 60345, A
835	96	5.7	640	4	US-09-906-618-292	Sequence 292, App	908	94	5.6	261	4	US-09-270-767-32898	Sequence 32898, A
836	96	5.7	647	3	US-08-753-007A-32	Sequence 32, Appl	909	94	5.6	261	4	US-09-270-767-48115	Sequence 48115, A
837	96	5.7	647	3	US-09-398-496-32	Sequence 32, Appl	910	94	5.6	422	1	US-08-036-555B-170	Sequence 170, App
838	96	5.7	650	3	US-09-310-463-2	Sequence 2, Appli	911	94	5.6	422	1	US-08-469-569-170	Sequence 3, Appli
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840	96	5.7	651	3	US-08-985-950-22	Sequence 22, App	913	94	5.6	422	1	US-08-249-322A-170	Sequence 170, App
841	96	5.7	651	4	US-09-546-049-22	Sequence 22, App	914	94	5.6	422	1	US-08-428-927-3	Sequence 3, Appli
842	96	5.7	653	4	US-08-467-602-212	Sequence 12, App	915	94	5.6	422	1	US-08-428-298-3	Sequence 3, Appli
843	96	5.7	653	4	US-08-411-295F-138	Sequence 138, App	916	94	5.6	422	1	US-08-339-517-3	Sequence 3, Appli
844	96	5.7	669	1	US-07-847-743B-8	Sequence 8, Appli	917	94	5.6	422	1	US-08-469-526A-170	Sequence 170, App
845	96	5.7	669	1	US-07-847-743B-13	Sequence 13, Appl	918	94	5.6	422	2	US-08-734-591A-170	Sequence 170, App
846	96	5.7	669	1	US-08-456-201-8	Sequence 8, Appli	919	94	5.6	422	2	US-08-469-660-170	Sequence 170, App
847	96	5.7	669	1	US-08-456-201-13	Sequence 13, Appl	920	94	5.6	422	2	US-08-341-018-72	Sequence 72, Appl
848	96	5.7	669	2	US-08-330-161-11	Sequence 11, Appl	921	94	5.6	422	3	US-08-470-335-170	Sequence 170, App
849	96	5.7	669	2	US-08-456-241-8	Sequence 8, Appli	922	94	5.6	422	3	US-08-735-021-170	Sequence 170, App
850	96	5.7	669	2	US-08-456-241-13	Sequence 13, Appl	923	94	5.6	422	3	US-08-734-664A-170	Sequence 170, App
851	96	5.7	669	2	US-08-440-401-11	Sequence 11, Appl	924	94	5.6	422	3	US-08-470-339-170	Sequence 170, App
852	96	5.7	669	2	US-08-419-878B-11	Sequence 11, Appl	925	94	5.6	422	4	US-08-467-602-170	Sequence 170, App
853	96	5.7	669	3	US-09-173-480-11	Sequence 11, Appl	926	94	5.6	422	4	US-08-411-295F-65	Sequence 65, Appl
854	96	5.7	669	5	PCT-US92-04295A-8	Sequence 8, Appli	927	94	5.6	422	4	US-08-411-295F-66	Sequence 66, Appl
855	96	5.7	669	5	PCT-US92-04295A-13	Sequence 13, Appl	928	94	5.6	422	4	US-08-411-295F-69	Sequence 69, Appl
856	96	5.7	687	4	US-08-467-602-254	Sequence 254, App	929	94	5.6	422	5	PCT-US94-05083C-166	Sequence 166, App
857	96	5.7	687	4	US-08-411-295F-180	Sequence 180, App	930	94	5.6	422	5	PCT-US94-05083C-185	Sequence 185, App
858	96	5.7	1140	4	US-09-579-692B-8	Sequence 8, Appli	931	94	5.6	422	5	PCT-US95-06846A-170	Sequence 170, App
859	96	5.7	1638	4	US-09-071-035-258	Sequence 258, App	932	94	5.6	458	6	5223394-7	Patent No. 5223394
860	96	5.7	1638	4	US-09-071-035-262	Sequence 262, App	933	94	5.6	458	6	5223394-7	Patent No. 5223394
861	96	5.7	1638	4	US-09-071-035-266	Sequence 266, App	934	94	5.6	635	3	US-08-470-335-247	Sequence 247, App
862	96	5.7	1747	4	US-09-134-000C-5999	Sequence 5999, App	935	94	5.6	635	3	US-08-467-602-302	Sequence 302, App
863	95.5	5.7	256	4	US-09-949-016-7326	Sequence 7326, Ap	936	94	5.6	635	4	US-08-411-295F-228	Sequence 228, App
864	95.5	5.7	652	3	US-09-310-463-4	Sequence 4, Appli	937	94	5.6	637	4	US-09-569-611C-35	Sequence 35, Appl
865	95.5	5.7	652	4	US-08-842-248A-4	Sequence 4, Appli	938	94	5.6	669	4	US-08-467-602-344	Sequence 344, App
866	95	5.6	287	3	US-08-862-124-17	Sequence 17, Appl	939	94	5.6	669	4	US-08-411-295F-270	Sequence 270, App
867	95	5.6	649	4	US-09-499-522-14	Sequence 14, Appl	940	94	5.6	852	3	US-08-985-526-34	Sequence 34, Appl
868	95	5.6	649	4	US-09-269-939A-8	Sequence 8, Appli	941	94	5.6	852	3	US-08-470-335-248	Sequence 248, App
869	95	5.6	879	1	US-08-554-612C-1	Sequence 1, Appli	942	94	5.6	852	4	US-08-467-602-300	Sequence 300, App
870	95	5.6	1501	2	US-08-447-464-3	Sequence 3, Appli	943	94	5.6	852	4	US-08-411-295F-226	Sequence 226, App
871	95	5.6	1501	2	US-08-716-679-3	Sequence 3, Appli	944	94	5.6	886	4	US-08-467-602-342	Sequence 342, App
872	94.5	5.6	194	3	US-08-630-172-14	Sequence 14, Appl	945	94	5.6	886	4	US-08-411-295F-268	Sequence 268, App
873	94.5	5.6	194	3	US-09-375-419-14	Sequence 14, Appl	946	94	5.6	899	3	US-08-470-335-249	Sequence 249, App
874	94.5	5.6	328	3	US-09-560-639-9	Sequence 9, Appli	947	94	5.6	899	3	US-08-467-602-301	Sequence 301, App
875	94.5	5.6	328	3	US-09-173-151A-25	Sequence 25, Appl	948	94	5.6	899	4	US-08-411-295F-227	Sequence 227, App
876	94.5	5.6	328	4	US-09-032-337-45	Sequence 45, Appl	949	94	5.6	933	4	US-08-467-602-343	Sequence 343, App
877	94.5	5.6	328	4	US-09-949-016-6424	Sequence 6424, Ap	950	94	5.6	933	4	US-08-411-295F-269	Sequence 269, App
878	94.5	5.6	477	2	US-08-359-705B-4	Sequence 4, Appli	951	94	5.6	983	3	US-09-412-554A-2	Sequence 2, Appli
879	94.5	5.6	477	2	US-08-286-846A-4	Sequence 4, Appli	952	94	5.6	143	2	US-08-653-402B-8	Sequence 8, Appli
880	94.5	5.6	477	2	US-08-457-880A-4	Sequence 4, Appli	953	93.5	5.5	259	3	US-09-560-639-11	Sequence 11, Appl
881	94.5	5.6	477	3	US-08-444-622A-4	Sequence 4, Appli	954	93.5	5.5	296	4	US-09-667-135-36	Sequence 36, Appl
882	94.5	5.6	477	3	US-08-942-562-4	Sequence 4, Appli	955	93.5	5.5	323	4	US-09-651-200-21	Sequence 21, Appl
883	94.5	5.6	477	3	US-09-156-923-4	Sequence 4, Appli	956	93.5	5.5	323	4	US-09-441-411-22	Sequence 22, Appl
884	94.5	5.6	483	4	US-09-949-016-8574	Sequence 8574, Ap	957	93.5	5.5	329	5	PCT-US94-09642-2	Sequence 2, Appli
885	94.5	5.6	556	3	US-09-560-639-8	Sequence 8, Appli	958	93.5	5.5	329	5	US-08-456-104-2	Sequence 2, Appli
886	94.5	5.6	610	2	US-08-724-394A-5	Sequence 5, Appli	959	93.5	5.5	329	2	US-08-101-624-2	Sequence 2, Appli
887	94.5	5.6	822	2	US-08-359-705B-2	Sequence 2, Appli	960	93.5	5.5	329	3	US-08-479-744A-2	Sequence 2, Appli
888	94.5	5.6	822	2	US-08-286-846A-2	Sequence 2, Appli	961	93.5	5.5	329	3	US-08-280-757B-2	Sequence 2, Appli
889	94.5	5.6	822	2	US-08-457-880A-2	Sequence 2, Appli	962	93.5	5.5	329	3	US-08-205-657A-23	Sequence 23, Appl
890	94.5	5.6	822	3	US-08-444-622A-2	Sequence 2, Appli	963	93.5	5.5	329	3	US-08-702-525-23	Sequence 23, Appl
891	94.5	5.6	822	3	US-08-542-562-2	Sequence 2, Appli	964	93.5	5.5	329	3	US-08-403-253A-4	Sequence 4, Appli
892	94.5	5.6	822	3	US-09-156-923-2	Sequence 2, Appli	965	93.5	5.5	329	4	US-08-435-816A-4	Sequence 4, Appli
893	94.5	5.6	822	4	US-09-949-016-6698	Sequence 6698, Ap	966	93.5	5.5	329	4	US-09-425-762-2	Sequence 2, Appli
894	94.5	5.6	847	1	US-08-286-305A-5	Sequence 5, Appli	967	93.5	5.5	329	4	US-09-837-867A-23	Sequence 23, Appl
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896	94.5	5.6	847	2	US-08-440-816A-5	Sequence 5, Appli	969	93.5	5.5	329	4	US-09-425-516-26	Sequence 26, Appl
897	94.5	5.6	847	3	US-09-417-381A-5	Sequence 35, Appl	970	93.5	5.5	329	4	US-09-425-516-26	Sequence 2, Appli
898	94.5	5.6	888	1	US-08-445-640-35	Sequence 35, Appl	971	93.5	5.5	329	5	PCT-US95-02576-23	Sequence 23, Appl
899	94.5	5.6	888	3	US-08-170-558-35	Sequence 35, Appl	972	93.5	5.5	372	4	US-09-949-016-11132	Sequence 11132, A
900	94.5	5.6	888	3	US-08-447-314-35	Sequence 35, Appl	973	93.5	5.5	388	1	US-08-445-640-12	Sequence 12, Appl
901	94.5	5.6	888	3	US-08-445-461-35	Sequence 35, Appl	974	93.5	5.5	388	3	US-08-170-558-12	Sequence 12, Appl
902	94.5	5.6	888	4	US-09-223-490-35	Sequence 35, Appl	975	93.5	5.5	388	3	US-08-447-314-12	Sequence 12, Appl
903	94.5	5.6	910	4	US-09-313-942-28	Sequence 28, Appl	976	93.5	5.5				

977	93.5	5.5	388	3	US-08-445-461-12	Sequence 12, Appl	1050	92	5.5	459	4	US-08-411-295F-225	Sequence 225, App
978	93.5	5.5	388	4	US-09-223-490-12	Sequence 12, Appl	1051	92	5.5	459	4	US-08-411-295F-288	Sequence 288, App
979	93.5	5.5	434	1	US-08-236-311-4	Sequence 4, Appli	1052	92	5.5	479	4	US-08-467-602-307	Sequence 307, App
980	93.5	5.5	434	3	US-08-457-918-4	Sequence 4, Appli	1053	92	5.5	479	4	US-08-411-295F-233	Sequence 233, App
981	93.5	5.5	434	4	US-10-157-408-4	Sequence 4, Appli	1054	92	5.5	490	4	US-08-467-602-345	Sequence 345, App
982	93.5	5.5	445	4	US-08-467-602-328	Sequence 328, App	1055	92	5.5	490	4	US-08-411-295F-271	Sequence 271, App
983	93.5	5.5	445	4	US-08-411-295F-254	Sequence 254, App	1056	92	5.5	493	4	US-08-467-602-341	Sequence 341, App
984	93.5	5.5	458	4	US-09-773-877B-26	Sequence 26, Appl	1057	92	5.5	493	4	US-08-411-295F-267	Sequence 267, App
985	93.5	5.5	479	4	US-08-467-602-370	Sequence 370, App	1058	92	5.5	513	4	US-08-467-602-349	Sequence 349, App
986	93.5	5.5	479	4	US-08-411-295F-296	Sequence 296, App	1059	92	5.5	513	4	US-08-411-295F-275	Sequence 275, App
987	93.5	5.5	770	1	US-08-525-654A-1	Sequence 1, Appli	1060	92	5.5	613	4	US-09-800-729-82	Sequence 82, Appl
988	93.5	5.5	771	1	US-08-525-654A-3	Sequence 3, Appli	1061	92	5.5	613	4	US-09-800-729-98	Sequence 98, Appl
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990	93	5.5	309	4	US-09-910-174B-7	Sequence 7, Appli	1063	92	5.5	644	4	US-08-467-602-311	Sequence 311, App
991	93	5.5	309	4	US-09-620-461-7	Sequence 7, Appli	1064	92	5.5	644	4	US-08-411-295F-237	Sequence 237, App
992	93	5.5	365	4	US-09-549-016-6907	Sequence 6907, Ap	1065	92	5.5	678	4	US-08-467-602-353	Sequence 353, App
993	93	5.5	391	4	US-09-949-016-7125	Sequence 7325, Ap	1066	92	5.5	678	4	US-08-411-295F-279	Sequence 279, App
994	93	5.5	523	4	US-09-910-174B-11	Sequence 11, Appl	1067	92	5.5	861	3	US-08-470-335-251	Sequence 251, App
995	93	5.5	523	4	US-09-620-461-11	Sequence 11, Appl	1068	92	5.5	861	4	US-08-467-602-312	Sequence 312, App
996	93	5.5	558	4	US-09-667-135-31	Sequence 31, Appl	1069	92	5.5	861	4	US-08-411-295F-238	Sequence 238, App
997	93	5.5	581	2	US-08-724-394A-3	Sequence 3, Appli	1070	92	5.5	895	4	US-08-467-602-354	Sequence 354, App
998	92.5	5.5	282	4	US-09-404-879A-393	Sequence 393, App	1071	92	5.5	895	4	US-08-411-295F-280	Sequence 280, App
999	92.5	5.5	282	4	US-09-667-857-393	Sequence 393, App	1072	92	5.5	908	3	US-08-470-335-252	Sequence 252, App
1000	92.5	5.5	309	4	US-09-404-879A-392	Sequence 392, App	1073	92	5.5	908	4	US-08-467-602-313	Sequence 313, App
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1002	92.5	5.5	329	4	US-09-667-135-32	Sequence 32, Appli	1075	92	5.5	942	4	US-08-467-602-355	Sequence 355, App
1003	92.5	5.5	389	4	US-08-467-602-276	Sequence 276, App	1076	92	5.5	942	4	US-08-411-295F-281	Sequence 281, App
1004	92.5	5.5	389	4	US-08-411-295F-202	Sequence 202, App	1077	91.5	5.4	272	4	US-09-728-219A-183	Sequence 183, App
1005	92.5	5.5	424	6	5169835-6	Patent No. 5169835	1078	91.5	5.4	272	4	US-09-318-786-37	Sequence 37, Appl
1006	92.5	5.5	424	6	5169835-6	Patent No. 5169835	1079	91.5	5.4	338	1	US-08-442-043A-17	Sequence 17, Appl
1007	92.5	5.5	426	1	US-08-336-583-2	Sequence 2, Appli	1080	91.5	5.4	338	3	US-08-441-893A-17	Sequence 17, Appl
1008	92.5	5.5	426	5	PCT-US95-13795-2	Sequence 2, Appli	1081	91.5	5.4	388	3	US-09-188-930-275	Sequence 275, App
1009	92.5	5.5	606	4	US-08-467-602-277	Sequence 277, App	1082	91.5	5.4	388	4	US-09-312-283C-275	Sequence 275, App
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1011	92.5	5.5	608	3	US-09-095-385-4	Sequence 4, Appli	1084	91.5	5.4	576	2	US-08-381-603-4	Sequence 4, Appli
1012	92.5	5.5	653	4	US-08-467-602-275	Sequence 275, App	1085	91.5	5.4	576	3	US-08-924-376-4	Sequence 4, Appli
1013	92.5	5.5	653	4	US-08-411-295F-201	Sequence 201, App	1086	91.5	5.4	576	3	US-08-685-212-4	Sequence 4, Appli
1014	92.5	5.5	746	3	US-08-434-000A-4	Sequence 4, Appli	1087	91.5	5.4	576	3	US-09-173-151A-30	Sequence 30, Appli
1015	92.5	5.5	746	3	US-09-312-157-4	Sequence 4, Appli	1088	91.5	5.4	576	4	US-08-466-932A-4	Sequence 4, Appli
1016	92.5	5.5	746	4	US-09-717-888-4	Sequence 4, Appli	1089	91.5	5.4	576	5	PCT-US94-02414-4	Sequence 4, Appli
1017	92.5	5.5	764	4	US-09-949-016-6254	Sequence 6254, Ap	1090	91.5	5.4	576	5	PCT-US96-08899-4	Sequence 4, Appli
1018	92.5	5.5	816	4	US-09-949-016-8119	Sequence 8119, Ap	1091	91.5	5.4	643	5	PCT-US93-00031-19	Sequence 19, Appl
1019	92.5	5.5	887	4	US-09-540-236-2911	Sequence 2911, Ap	1092	91.5	5.4	644	5	PCT-US93-00031-21	Sequence 21, Appl
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1021	92	5.5	274	4	US-09-813-659-30	Sequence 30, Appl	1094	91.5	5.4	647	3	US-08-482-073-5	Sequence 5, Appli
1022	92	5.5	274	4	US-09-549-067A-30	Sequence 30, Appl	1095	91.5	5.4	647	5	PCT-US93-00031-11	Sequence 11, Appl
1023	92	5.5	302	1	US-08-121-054C-18	Sequence 18, Appl	1096	91.5	5.4	647	5	PCT-US93-00031-23	Sequence 23, Appl
1024	92	5.5	302	1	US-08-121-054C-30	Sequence 30, Appl	1097	91.5	5.4	740	5	PCT-US93-00031-17	Sequence 17, Appl
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1034	92	5.5	414	3	US-08-470-339-188	Sequence 188, App	1107	91	5.4	405	4	US-08-467-602-384	Sequence 384, App
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1038	92	5.5	422	4	US-08-411-295F-250	Sequence 250, App	1111	91	5.4	526	1	US-08-471-570-4	Sequence 4, Appli
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1040	92	5.5	425	4	US-08-467-602-320	Sequence 320, App	1113	91	5.4	572	4	US-08-411-295F-161	Sequence 161, App
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1044	92	5.5	456	4	US-08-467-602-366	Sequence 366, App	1117	91	5.4	619	4	US-08-411-295F-159	Sequence 159, App
1045	92	5.5	456	4	US-08-411-295F-229	Sequence 229, App	1118	91	5.4	652	1	US-08-471-570-10	Sequence 10, Appl
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1048	92	5.5	459	4	US-08-467-602-299	Sequence 299, App	1121	90.5	5.4	238	4	US-09-798-689-21	Sequence 21, Appl
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1174	90	5.3	659	4	US-08-411-295F-155	Sequence 155, App	1247	87.5	5.2	684	4	US-08-411-295F-185	Sequence 185, App
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1189	89	5.3	287	4	US-09-800-729-153	Sequence 153, App	1262	86.5	5.1	395	4	US-08-467-602-290	Sequence 216, App
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1191	89	5.3	342	4	US-09-684-708A-27	Sequence 27, Appli	1264	86.5	5.1	612	4	US-08-467-602-291	Sequence 217, App
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1273	86.5	5.1	894	1	US-08-445-640-34	Sequence 34, Appl1	1346	85	5.0	361	4	US-08-411-895F-174	Sequence 174, App
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1276	86.5	5.1	894	3	US-08-445-461-34	Sequence 34, Appl1	1349	85	5.0	578	4	US-08-411-295F-175	Sequence 175, App
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1278	86.5	5.1	904	4	US-09-877-730-6	Sequence 6, Appl1	1351	85	5.0	625	4	US-08-467-602-250	Sequence 250, App
1279	86.5	5.1	907	4	US-09-877-730-20	Sequence 20, Appl1	1352	85	5.0	625	4	US-08-411-295F-176	Sequence 176, App
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1281	86.5	5.1	985	4	US-09-877-730-10	Sequence 10, Appl1	1354	85	5.0	769	3	US-09-312-157-10	Sequence 10, Appl
1282	86.5	5.1	991	4	US-09-877-730-12	Sequence 12, Appl1	1355	85	5.0	769	4	US-09-717-888-10	Sequence 10, Appl
1283	86.5	5.1	1018	1	US-08-452-052-2	Sequence 2, Appl1	1356	85	5.0	2409	6	5180808-2	Patent No. 5180808
1284	86.5	5.1	1069	4	US-09-877-730-2	Sequence 2, Appl1	1357	85	5.0	2409	6	5180808-2	Patent No. 5180808
1285	86.5	5.1	1070	4	US-09-877-730-18	Sequence 18, Appl1	1358	84.5	5.0	119	4	US-09-858-664A-30	Sequence 30, Appl1
1286	86.5	5.1	1152	4	US-09-877-730-8	Sequence 8, Appl1	1359	84.5	5.0	119	4	US-10-274-978-31	Sequence 31, Appl1
1287	86.5	5.1	1163	4	US-09-375-248-19	Sequence 19, Appl1	1360	84.5	5.0	119	4	US-10-697-263-31	Sequence 31, Appl1
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1289	86	5.1	241	3	US-08-470-335-195	Sequence 195, App	1362	84.5	5.0	249	2	US-08-797-689-18	Sequence 18, Appl1
1290	86	5.1	241	3	US-08-470-339-195	Sequence 195, App	1363	84.5	5.0	249	4	US-09-984-186-18	Sequence 18, Appl1
1291	86	5.1	241	3	US-08-467-602-389	Sequence 389, App	1364	84.5	5.0	419	6	5169835-2	Patent No. 5169835
1292	86	5.1	241	4	US-08-411-295F-47	Sequence 47, Appl1	1365	84.5	5.0	419	6	5169835-2	Patent No. 5169835
1293	86	5.1	260	4	US-09-949-016-8243	Sequence 8243, Ap	1366	84.5	5.0	457	1	US-08-416-478A-8	Sequence 8, Appl1
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1295	86	5.1	398	1	US-08-091-519-2	Sequence 2, Appl1	1368	84.5	5.0	457	2	US-08-394-442B-8	Sequence 8, Appl1
1296	86	5.1	398	1	US-08-442-043A-2	Sequence 2, Appl1	1369	84.5	5.0	524	4	US-09-270-767-44009	Sequence 44009, A
1297	86	5.1	398	3	US-09-173-151A-26	Sequence 26, Appl1	1370	84	5.0	524	4	US-09-471-276-855	Sequence 855, App
1298	86	5.1	398	4	US-09-461-908-2	Sequence 2, Appl1	1371	84	5.0	122	3	US-08-811-682-8	Sequence 8, Appl1
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1301	86	5.1	398	4	US-09-321-667-8	Sequence 8, Appl1	1374	84	5.0	241	2	US-08-224-591-18	Sequence 18, Appl1
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1323	85.5	5.1	403	4	US-09-638-648-5	Sequence 5, Appl1	1396	84	5.0	603	4	US-08-411-295F-205	Sequence 205, App
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1325	85.5	5.1	477	4	US-09-949-016-9193	Sequence 9193, Ap	1398	84	5.0	650	4	US-08-467-602-206	Sequence 206, App
1326	85.5	5.1	477	4	US-09-949-016-9194	Sequence 9194, Ap	1399	84	5.0	650	4	US-08-858-664A-25	Sequence 25, Appl
1327	85.5	5.1	477	4	US-09-949-016-9195	Sequence 9195, Ap	1400	83.5	4.9	100	4	US-08-467-602-26	Sequence 26, Appl1
1328	85.5	5.1	477	4	US-09-949-016-9196	Sequence 9196, Ap	1401	83.5	4.9	100	4	US-10-274-978-26	Sequence 26, Appl1
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1330	85.5	5.1	477	4	US-09-949-016-9198	Sequence 9198, Ap	1403	83.5	4.9	231	1	US-08-168-091A-4	Sequence 4, Appl1
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1334	85.5	5.1	741	4	US-09-585-858-11	Sequence 11, Appl1	1407	83.5	4.9	288	2	US-08-456-104-6	Sequence 6, Appl1
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1341	85	5.0	247	4	US-09-949-016-6225	Sequence 6225, Ap	1414	83.5	4.9	288	3	US-08-205-697A-19	Sequence 19, Appl1

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RESULT 2
US-09-953-499-2
; Sequence 2, Application US/09953499
; Patent No. 6838554
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Fong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Gurney, Austin L.
; APPLICANT: Napier, Mary A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: COMPOUNDS, COMPOSITIONS AND METHODS FOR THE TREATMENT
; OF DISEASES CHARACTERIZED BY A33- RELATED ANTIGENS
; FILE REFERENCE: P1216R1 (US)
; CURRENT APPLICATION NUMBER: US/09/953,499
; CURRENT FILING DATE: 2001-09-14
; PRIOR APPLICATION NUMBER: US/09/254,465
; PRIOR FILING DATE: 1999-03-05
; PRIOR APPLICATION NUMBER: PCT/US98/24855
; PRIOR FILING DATE: 1998-11-20
; PRIOR APPLICATION NUMBER: US 60/066,364
; PRIOR FILING DATE: 1997-11-21
; PRIOR APPLICATION NUMBER: US 60/078,936
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: PCT/US98/19437
; PRIOR FILING DATE: 1998-09-17
; NUMBER OF SEQ ID NOS: 30
; SEQ ID NO 2
; LENGTH: 321
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-953-499-2

Query Match 100.0%; Score 1688; DB 4; Length 321;
Best Local Similarity 100.0%; Pred. No. 1.4e-165;
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy 121 DGNQVVRDKITELRVKLSVSKPTVTTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180
Db 121 DGNQVVRDKITELRVKLSVSKPTVTTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180

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Qy 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300

Qy 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAAR 321
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RESULT 3
US-09-369-247-63
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; Sequence 63, Application US/09369247
; Patent No. 656992
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 44 Human Secreted Proteins
; FILE REFERENCE: P2024P1
; CURRENT APPLICATION NUMBER: US/09/369,247
; CURRENT FILING DATE: 1999-08-05
; EARLIER APPLICATION NUMBER: 60/074,118
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,157
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,137
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,341
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,141
; EARLIER FILING DATE: 1998-02-09
; NUMBER OF SEQ ID NOS: 172
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 63
; LENGTH: 306
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (306)
; OTHER INFORMATION: Xaa equals stop translation
US-09-369-247-63

Query Match 67.4%; Score 1137; DB 4; Length 306;
Best Local Similarity 70.4%; Pred. No. 9.4e-109;
Matches 226; Conservative 0; Mismatches 1; Indels 94; Gaps 1;

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Qy 121 DGNQVVRDKITELRVKLSVSKPTVTTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180
Db 121 DGNQVVRDKITELRVQ----- 136

Qy 181 QTNNOEPIKVATLSTLLFPKPAVIADSGSYFCTAKGVGSEQHSDIVKFFVKDSSKLLKTK 240
Db 137 -----KHSSKLLKTK 146

Qy 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300
Db 147 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 206

Qy 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 207 MAYIMLCRKTSQOEHVYEAAAR 227

RESULT 4
US-09-763-902B-6
; Sequence 6, Application US/09763902B
; Patent No. 6838241
; GENERAL INFORMATION:
; APPLICANT: INCYTE PHARMACEUTICALS, INC.
; APPLICANT: TANG, Y. Tom
; APPLICANT: LAL, Preeti
; APPLICANT: BANDMAN, Olga
; APPLICANT: YUE, Henry
; APPLICANT: CORLEY, Neil C.
; APPLICANT: GUEGLER, Karl J.
; APPLICANT: GORGONE, Gina A.
; APPLICANT: BAUGHN, Mariah R.
```

APPLICANT: PATTERSON, Chandra
TITLE OF INVENTION: PROTEIN TRANSPORT-ASSOCIATED MOLECULES
FILE REFERENCE: PF-0577 PCT
CURRENT APPLICATION NUMBER: US/09/763,902B
CURRENT FILING DATE: 2002-08-22
PRIOR APPLICATION NUMBER: 60/098,206
PRIOR FILING DATE: 1998-08-27
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PERL Program
SEQ ID NO 6
LENGTH: 175
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. 68382411871275CD1
US-09-763-902B-6

Query Match 51.5%; Score 870; DB 4; Length 175;
Best Local Similarity 64.7%; Pred. No. 1.4e-81;
Matches 174; Conservative 0; Mismatches 1; Indels 94; Gaps 1;
QY 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGDNLPCTYDPLQGYTQVLVKLVQR 60
DB 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGDNLPCTYDPLQGYTQVLVKLVQR 60
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLSTLEMDRSHYTCVWTQTP 120
DB 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLSTLEMDRSHYTCVWTQTP 120
QY 121 DGNQVVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 180
DB 121 DGNQVVRDKITELRVQ- 136
QY 181 QTNQPEIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEQHSDIVKVVVDKSSKLLTKT 240
DB 137 -----KHSSKLLTKT 146
QY 241 TEAPTTMTYPLKATSVKQSWDWTMDMG 269
DB 147 TEAPTTMTYPLKATSVKQSWDWTMDMG 175

RESULT 5
US-09-188-930-189
Sequence 189, Application US/09188930A
Patent No. 6150502
GENERAL INFORMATION:
APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 189
LENGTH: 299
TYPE: PRT
ORGANISM: Human
FEATURE:
NAME/KEY: UNSURE
LOCATION: (247)...(247)
NAME/KEY: UNSURE
LOCATION: (289)...(289)
US-09-188-930-189

Query Match 10.6%; Score 178.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;

Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;
QY 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGDNLPCTYDPLQGYTQVLVKLVQR 60
DB 17 LAILLCSLALGSVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW--- 62
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHK-----VPGDVSLSLSTLEMDRSHYTCV 115
DB 63 -----KFDQGDVTRLVVCYNNKITASYEDRVTFPLGTGTFKSVTRE--DTGTVTCMV 111
QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 174
DB 112 SEEGNSYGEVKKLVIL-----VPPSKPTVNIPS-----SATIGNRAVLTCSDQSGSPSE 163
QY 175 YIWK-----QQTN-----NQEPIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEQH 222
DB 164 YTFKDGIVMPTNPKSTRAFSNSSVYLNPTTGGELVFDPLSLASDTGECSEARNGYGTPT 223
QY 223 SDIVK 227
DB 224 SNAVR 228

RESULT 6
US-09-188-930-331
Sequence 331, Application US/09188930A
Patent No. 6150502
GENERAL INFORMATION:
APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Murison, James Greg
TITLE OF INVENTION: Compositions Isolated From Skin Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 331
LENGTH: 299
TYPE: PRT
ORGANISM: Human
US-09-188-930-331

Query Match 10.6%; Score 178.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;
QY 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGDNLPCTYDPLQGYTQVLVKLVQR 60
DB 17 LAILLCSLALGSVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW--- 62
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHK-----VPGDVSLSLSTLEMDRSHYTCV 115
DB 63 -----KFDQGDVTRLVVCYNNKITASYEDRVTFPLGTGTFKSVTRE--DTGTVTCMV 111
QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 174
DB 112 SEEGNSYGEVKKLVIL-----VPPSKPTVNIPS-----SATIGNRAVLTCSDQSGSPSE 163
QY 175 YIWK-----QQTN-----NQEPIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEQH 222
DB 164 YTFKDGIVMPTNPKSTRAFSNSSVYLNPTTGGELVFDPLSLASDTGECSEARNGYGTPT 223
QY 223 SDIVK 227
DB 224 SNAVR 228

RESULT 7
US-09-462-270-2

Qy	223	SDIVK	227
		:	:
Db	224	SNAVR	228

RESULT 10
US-09-312-283C-331
; Sequence 331, Application US/09312283C

```

/ GENERATED INFORMATION:
/ APPLICANT: Watson, James D.
/ APPLICANT: Strachan, Lorna
/ APPLICANT: Sleeman, Matthew
/ APPLICANT: Onrust, Rene
/ APPLICANT: Murison, James G.
/ APPLICANT: Kumble, Krishanand D.
/ TITLE OF INVENTION: Compositions Isolated from Skin Cells
/ TITLE OF INVENTION: and Methods for Their Use
/ FILE REFERENCE: 11000.1011c2
/ CURRENT APPLICATION NUMBER: US/09/312,283C
/ CURRENT FILING DATE: 1999-05-14
/ NUMBER OF SEQ ID NOS: 425
/ SOFTWARE: FastSeq for Windows Version 4.0
/

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Query Match	10.6%	Score 178.5;	DB 4;	Length 299;
Best Local Similarity	26.5%;	Pred. No. 7.8e-10;		
Matches 65: Conservative	36;	Mismatches 93;	Indels 51;	Gaps 11;

Qy	60	1	MGILLGILLLLGHLTVDYGRPTILEVPESVTFGWGDVNLPCCTYPDQGYQVLVKWLQVR	60
Db	62	17	LAILLCSLALGVSPTVHHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW----	62
Qy	115	61	GSDPVTIFURDSSGDHIOQAQKQGRHLVSHK-----VPGDVSLQLSFTLEMDRSHYTCEV	115
Db	111	63	-----KFOGDTTRLVCYNNKITASYEDRVTFELPTGITPKSVTRE--DTGYTCMV	111
Qy	174	116	TWQTPDGNQVRDKTEILRVQKLSVSKPTVTGSGGYFTVPQGNRISLQCOAR-GSPPTS	174
Db	163	112	SBEGGNSYGEVVKLIVL-----VPPSKPTNIPS-----SATIGNRAVLTCSEQDGGPPSE	163
Qy	222	175	YTWYK-----QQTN-----NQBEPIKVATLSTLLEKPAVIADSGSYFCTAKGVGSEQH	222
Db	223	164	YTWFKDGIWNPINPKSTRAFNSSSYVLNPTTGGELVFDPLSASDTGEYSCBARNGYTGMT	223
Qy	227	223	SDIVK	227
Db	228	224	SNAVR	228

RESULT 11
US-09-907-794A-119
: Sequence 119. Application US/09907794A

APPLICANT:	Genentech, Inc.
APPLICANT:	Askenazi, Avi
APPLICANT:	Botstein, David
APPLICANT:	Desnovers, Luc
APPLICANT:	Eaton, Dan L.
APPLICANT:	Ferrara, Napoleone
APPLICANT:	Filvaroff, Ellen
APPLICANT:	Fong, Sherman
APPLICANT:	Gao, Wei-Qiang
APPLICANT:	Gerbet, Hanspeter
APPLICANT:	Gerritsen, Mary E.
APPLICANT:	Goddard, A.
APPLICANT:	Godowski, Paul J.
APPLICANT:	Grimaldi, Christopher

? APPLICANT: Gurney, Austin L.
 ? APPLICANT: Hillan, Kenneth, J.
 ? APPLICANT: Kljavin, Ivar J.
 ? APPLICANT: Mather, Jennie P.
 ? APPLICANT: Pan, James
 ? APPLICANT: Paoni, Nicholas F.
 ? APPLICANT: Roy, Margaret Ann
 ? APPLICANT: Stewart, Timothy A.
 ? APPLICANT: Tumas, Daniel
 ? APPLICANT: Williams, P. Mickey
 ? APPLICANT: Wood, William, I.
 ? TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 ?
 ? FILE OF INVENTION: Acids Encoding the Same
 ? FILE REFERENCE: 10466-14
 ? CURRENT APPLICATION NUMBER: US/09/907,794A

Query Match	10.6%	Score 178.5	DB 4	Length 299
Best Local Similarity	26.5%	Pred. No. 7.8e-10		
Matches 65	Conservative 36	Mismatches 93	Indels 51	Gaps 11
Qy	1	MGILLGLLLGHLTVDTGYGRPILEVEPSVTGPKKGDVNLCTVDPLQGYQTQVLKVLVQR	60	
Db	17	LAILLCSLAGSVTVHS-SEPEVRIPEN-----NPVKLSCAV---SGFSSPRVEW----	62	
Qy	61	GSDPVITFLDSSGDHIQAKYQCRHLVSHK-----VPGDVSLQLSTLEMDRSHYTCV	115	
Db	63	-----KFDQGDTRLVCYNNKITSASYEDRVTFPLTGITPKSVTRE--DGTGTYTCMV	111	
Qy	116	TWTPDGNQVVRDKITELRVOKLSVKPPTVTGSGYGVTFVQGNRISLQCOAR-GSPDIS	174	
Db	112	SEGGNSYGEVKKVLIIVL-----VPPSKPTVNIIPS-----SATIGNRAVLTCSEODGSPSE	163	

FILE REFERENCE: 011.00221
CURRENT APPLICATION NUMBER: US/09/397,243D
CURRENT FILING DATE: 1999-09-16
PRIOR APPLICATION NUMBER: 60/100,638
PRIOR FILING DATE: 1998-09-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patent in Ver. 2.1
SEQ ID NO 3
LENGTH: 299
TYPE: PRT
ORGANISM: Homo sapiens
US-09-397-243D-3

Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGIALLGLLHLTVDTYGRPILEVPESVTGPMKGDVNLPCYDPLQGYTQVLVKLVQR 60
Db 17 LAILLCSLALGSVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62

QY 61 GSDPVTIFLRDSSGDHIQQAQYGRHLVSHK-----VPGDVSLLQSLTLEMDRSHYTCV 115
Db 63 -----KFDQGDTRLLVCYNNKITASYEDRVTFPLTGITFKSVTRE--DTGTYTCMV 111

QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGFVTPQGMRLISLQCAR-GSPPTS 174
Db 112 SEEGGNSYGEVKVLIVL-----VPPSKPTVNIIPS-----SATIGNRAVLTCSEODGSPPE 163

QY 175 YIWK-----QQTN-----NOEPIKVAATLSTLFLKPAVIADSGSYFCTAGQGVSEQH 222
Db 164 YTFKDGIVMPTNPKSTRAFSNSYVNLPTTGELVDFPLSDTGEYSCEARNGYGTPTMT 223

QY 223 SDIVK 227
Db 224 SNAVR 228

US-09-902-775A-119

Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGIALLGLLHLTVDTYGRPILEVPESVTGPMKGDVNLPCYDPLQGYTQVLVKLVQR 60
Db 17 LAILLCSLALGSVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62

QY 61 GSDPVTIFLRDSSGDHIQQAQYGRHLVSHK-----VPGDVSLLQSLTLEMDRSHYTCV 115
Db 63 -----KFDQGDTRLLVCYNNKITASYEDRVTFPLTGITFKSVTRE--DTGTYTCMV 111

QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGFVTPQGMRLISLQCAR-GSPPTS 174
Db 112 SEEGGNSYGEVKVLIVL-----VPPSKPTVNIIPS-----SATIGNRAVLTCSEODGSPPE 163

QY 175 YIWK-----QQTN-----NOEPIKVAATLSTLFLKPAVIADSGSYFCTAGQGVSEQH 222
Db 164 YTFKDGIVMPTNPKSTRAFSNSYVNLPTTGELVDFPLSDTGEYSCEARNGYGTPTMT 223

QY 223 SDIVK 227
Db 224 SNAVR 228

RESULT 14
US-09-397-243D-3
Sequence 3, Application US/09397243D
Patent No. 6699688
GENERAL INFORMATION:
APPLICANT: Kornecki, Elizabeth
APPLICANT: Sobocka, Malgorzata B.
TITLE OF INVENTION: Human Platelet F11 Receptor

RESULT 15
US-09-906-700-119
Sequence 119, Application US/09906700
Patent No. 6723535
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kijavini, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.

TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
Acids Encoding the Same
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/906,700
CURRENT FILING DATE: 2000-09-18

691	178.5	10.6	299	20	US-11-009-868-1	Sequence 1, Appli	906	158	9.4	352	14	US-10-223-084-280	Sequence 280, App
692	178.5	10.6	320	16	US-10-128-558-176	Sequence 376, App	907	158	9.4	352	14	US-10-223-088-280	Sequence 280, App
693	178.5	10.6	336	16	US-10-128-558-193	Sequence 193, App	908	158	9.4	352	14	US-10-223-090-280	Sequence 280, App
694	177	10.5	319	9	US-09-981-353-65	Sequence 65, Appli	913	158	9.4	352	14	US-10-223-087-280	Sequence 280, App
695	177	10.5	319	9	US-09-953-499-6	Sequence 6, Appli	914	158	9.4	352	14	US-10-223-083-280	Sequence 280, App
696	177	10.5	319	14	US-10-265-542-6	Sequence 6, Appli	917	158	9.4	352	14	US-10-223-089-280	Sequence 280, App
697	177	10.5	319	15	US-10-295-027-1165	Sequence 1165, Ap	919	158	9.4	352	14	US-10-223-081-280	Sequence 280, App
698	177	10.5	319	16	US-10-633-008-6	Sequence 6, Appli	938	158	9.4	352	14	US-10-223-082-280	Sequence 280, App
699	177	10.5	319	16	US-10-785-220-6	Sequence 6, Appli	940	158	9.4	352	15	US-10-305-654-280	Sequence 280, App
700	177	10.5	319	16	US-10-785-221-6	Sequence 6, Appli	943	158	9.4	352	15	US-10-081-056-280	Sequence 280, App
701	177	10.5	319	16	US-10-785-433-6	Sequence 6, Appli	948	158	9.4	365	9	US-09-971-798-2	Sequence 2, Appli
702	177	10.5	319	16	US-10-767-374-6	Sequence 6, Appli	949	158	9.4	365	14	US-10-176-847-78	Sequence 78, Appl
703	177	10.5	319	16	US-10-785-607-6	Sequence 6, Appli	950	158	9.4	365	17	US-10-482-029-76	Sequence 76, Appl
704	177	10.5	319	16	US-10-785-607-6	Sequence 6, Appli	951	158	9.4	365	18	US-10-926-337A-3	Sequence 3, Appli
705	177	10.5	336	15	US-10-363-616-318	Sequence 318, App	952	158	9.4	505	14	US-10-114-153-12	Sequence 12, Appl
706	176	10.4	268	14	US-10-265-542-24	Sequence 24, Appl	953	157	9.3	323	9	US-09-971-798-31	Sequence 31, Appl
707	176	10.4	268	16	US-10-633-008-24	Sequence 24, Appl	954	157	9.3	343	9	US-09-971-798-27	Sequence 27, Appl
708	176	10.4	270	9	US-09-953-499-24	Sequence 24, Appl	955	153.5	9.1	373	9	US-09-796-858-24	Sequence 24, Appl
709	176	10.4	270	16	US-10-785-220-24	Sequence 24, Appl	987	153.5	9.1	373	10	US-09-997-428-503	Sequence 503, App
710	176	10.4	270	16	US-10-785-221-24	Sequence 24, Appl	1093	153.5	9.1	373	14	US-10-167-749-59	Sequence 59, Appl
711	176	10.4	270	16	US-10-785-433-24	Sequence 24, Appl	1245	153.5	9.1	373	14	US-10-223-085-64	Sequence 64, Appl
712	176	10.4	270	16	US-10-767-374-24	Sequence 24, Appl	1251	153.5	9.1	373	14	US-10-219-065-130	Sequence 130, App
713	176	10.4	270	16	US-10-785-607-24	Sequence 24, Appl	1280	153.5	9.1	373	14	US-10-223-084-64	Sequence 64, Appl
714	176	10.4	270	18	US-10-767-904-24	Sequence 24, Appl	1281	153.5	9.1	373	14	US-10-223-088-64	Sequence 64, Appl
715	176	10.4	273	9	US-09-953-499-26	Sequence 26, Appl	1282	153.5	9.1	373	14	US-10-223-090-64	Sequence 64, Appl
716	176	10.4	273	14	US-10-265-542-26	Sequence 26, Appl	1287	153.5	9.1	373	14	US-10-223-087-64	Sequence 64, Appl
717	176	10.4	273	16	US-10-633-008-26	Sequence 26, Appl	1289	153.5	9.1	373	14	US-10-223-083-64	Sequence 64, Appl
718	176	10.4	273	16	US-10-785-220-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
719	176	10.4	273	16	US-10-785-221-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
720	176	10.4	273	16	US-10-785-433-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
721	176	10.4	273	16	US-10-767-374-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
722	176	10.4	273	16	US-10-785-607-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
723	176	10.4	273	18	US-10-767-904-26	Sequence 26, Appl	1292	153.5	9.1	373	14	US-10-223-089-64	Sequence 64, Appl
724	172.5	10.2	316	16	US-10-785-351-13	Sequence 13, Appl	1463	153.5	9.1	373	14	US-10-223-081-64	Sequence 64, Appl
725	171	10.1	47	16	US-10-425-115-299346	Sequence 299346,							
726	169.5	10.0	260	9	US-09-953-499-23	Sequence 23, Appl							
727	169.5	10.0	260	14	US-10-265-542-23	Sequence 23, Appl							
728	169.5	10.0	260	16	US-10-633-008-23	Sequence 23, Appl							
729	169.5	10.0	260	16	US-10-785-220-23	Sequence 23, Appl							
730	169.5	10.0	260	16	US-10-785-221-23	Sequence 23, Appl							
731	169.5	10.0	260	16	US-10-785-433-23	Sequence 23, Appl							
732	169.5	10.0	260	16	US-10-767-374-23	Sequence 23, Appl							
733	169.5	10.0	260	16	US-10-785-607-23	Sequence 23, Appl							
734	169.5	10.0	260	18	US-10-767-904-23	Sequence 23, Appl							
735	169.5	10.0	263	9	US-09-953-499-25	Sequence 25, Appl							
736	169.5	10.0	263	14	US-10-265-542-25	Sequence 25, Appl							
737	169.5	10.0	263	16	US-10-633-008-25	Sequence 25, Appl							
738	169.5	10.0	263	16	US-10-785-220-25	Sequence 25, Appl							
739	169.5	10.0	263	16	US-10-785-221-25	Sequence 25, Appl							
740	169.5	10.0	263	16	US-10-785-433-25	Sequence 25, Appl							
741	169.5	10.0	263	16	US-10-767-374-25	Sequence 25, Appl							
742	169.5	10.0	263	16	US-10-785-607-25	Sequence 25, Appl							
743	169.5	10.0	263	18	US-10-767-904-25	Sequence 25, Appl							
744	167	9.9	300	9	US-09-953-499-10	Sequence 10, Appl							
745	167	9.9	300	14	US-10-265-542-10	Sequence 10, Appl							
746	167	9.9	300	16	US-10-633-008-10	Sequence 10, Appl							
747	167	9.9	300	16	US-10-785-220-10	Sequence 10, Appl							
748	167	9.9	300	16	US-10-785-221-10	Sequence 10, Appl							
749	167	9.9	300	16	US-10-785-433-10	Sequence 10, Appl							
750	167	9.9	300	16	US-10-767-374-10	Sequence 10, Appl							
751	167	9.9	300	16	US-10-785-607-10	Sequence 10, Appl							
752	167	9.9	300	16	US-10-785-351-12	Sequence 12, Appl							
753	167	9.9	300	18	US-10-767-904-10	Sequence 10, Appl							
754	166	9.8	365	9	US-09-899-634A-4	Sequence 4, Appli							
755	165.5	9.8	36946	18	US-10-840-512-155	Sequence 155, App							
756	163.5	9.7	261	9	US-09-899-634A-2	Sequence 2, Appli							
757	158	9.4	208	18	US-10-481-090A-8	Sequence 8, Appli							
784	158	9.4	352	10	US-09-997-428-505	Sequence 505, App							
829	158	9.4	352	13	US-10-053-107-10	Sequence 10, Appl							
835	158	9.4	352	14	US-10-213-145-10	Sequence 10, Appl							
843	158	9.4	352	14	US-10-213-199-10	Sequence 10, Appl							
874	158	9.4	352	14	US-10-223-085-280	Sequence 280, App							
880	158	9.4	352	14	US-10-219-065-216	Sequence 216, App							

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